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(54) Title: IL-17 RECEPTOR LIKE MOLECULES AND USES THEREOF

(57) Abstract: Novel IL-17 receptor like polypeptides and nucleic acid molecules encoding the same. The invention also pro-  
vides vectors, host cells, agonists and antagonists (including selective binding agents), and method for producing IL-17 receptor like  
polypeptides. Also provided for are methods for the treatment, diagnosis, amelioration, or prevention of diseases with IL-17 receptor  
like polypeptides.

IL-17 RECEPTOR LIKE MOLECULES AND USES THEREOFRelated Application

5       The present application claims priority from U.S. patent application 09/724,460 file November 28, 2000 which claims priority from provisional application US 60/189,816 filed March 16, 2000.

10   Field of the Invention

      The present invention relates to novel IL-17 receptor like polypeptides and nucleic acid molecules encoding the same. The invention also relates to vectors, host cells, pharmaceutical compositions, selective binding agents and  
15   methods for producing IL-17 receptor like polypeptides. Also provided for are methods for the diagnosis, treatment, amelioration, and/or prevention of diseases associated with IL-17 receptor like polypeptides.

20   Background of the Invention

      Technical advances in the identification, cloning, expression and manipulation of nucleic acids have greatly accelerated the discovery of novel therapeutics based upon deciphering the human genome. Rapid nucleic acid sequencing  
25   techniques can now generate sequence information at unprecedented rates and, coupled with computational analyses, allow the assembly of overlapping sequences into partial and entire genomes and the identification of polypeptide-encoding regions. A comparison of a predicted amino acid sequence  
30   against a database compilation of known amino acid sequences can allow one to determine the extent of homology to previously identified sequences and/or structural landmarks. The cloning and expression of a polypeptide-encoding region of

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a nucleic acid molecule provides a polypeptide product for structural and functional analyses. The manipulation of nucleic acid molecules and encoded polypeptides to create variants and derivatives thereof may confer advantageous properties on a product for use as a therapeutic.

In spite of the significant technical advances in genome research over the past decade, the potential for the development of novel therapeutics based on the human genome is still largely unrealized. Many genes encoding potentially beneficial polypeptide therapeutics, or those encoding polypeptides, which may act as "targets" for therapeutic molecules, have still not been identified. In addition, structural and functional analyses of polypeptide products from many human genes have not been undertaken.

Accordingly, it is an object of the invention to identify novel polypeptides and nucleic acid molecules encoding the same, which have diagnostic or therapeutic benefit.

#### Summary of the Invention

The present invention relates to novel IL-17 receptor like nucleic acid molecules and encoded polypeptides.

The invention provides for an isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:

(a) the nucleotide sequence as set forth in SEQ ID NO: 1;

(b) a nucleotide sequence encoding the polypeptide as set forth in SEQ ID NO: 2;

(c) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of (a) or (b), wherein the encoded polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

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(d) a nucleotide sequence complementary to any of (a)-(c).

5 The invention also provides for an isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:

(a) a nucleotide sequence encoding a polypeptide that is at least about 70, 75, 80, 85, 90, 95, 96, 97, 98, or 99  
10 percent identical to the polypeptide as set forth in SEQ ID NO: 2, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(b) a nucleotide sequence encoding an allelic variant or splice variant of the nucleotide sequence as set forth in SEQ  
15 ID NO: 1, wherein the encoded polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(c) a nucleotide sequence of SEQ ID NO: 1, (a), or (b) encoding a polypeptide fragment of at least about 25 amino acid residues, wherein the polypeptide has an activity of the  
20 polypeptide as set forth in SEQ ID NO: 2;

(d) a nucleotide sequence of SEQ ID NO: 1, or (a)-(c) comprising a fragment of at least about 16 nucleotides;

(e) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of  
25 any of (a)-(d), wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

(f) a nucleotide sequence complementary to any of (a)-(e).

30 The invention further provides for an isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:



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(a) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one conservative amino acid substitution, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

5 (b) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one amino acid insertion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

10 (c) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one amino acid deletion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

15 (d) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 which has a C- and/or N- terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

20 (e) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one modification selected from the group consisting of amino acid substitutions, amino acid insertions, amino acid deletions, C-terminal truncation, and N-terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(f) a nucleotide sequence of (a)-(e) comprising a fragment of at least about 16 nucleotides;

25 (g) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of any of (a)-(f), wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

30 (h) a nucleotide sequence complementary to any of (a)-(e).

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The invention also provides for an isolated polypeptide comprising the amino acid sequence selected from the group consisting of:

(a) an amino acid sequence for an ortholog of SEQ ID NO: 2, wherein the encoded polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(b) an amino acid sequence that is at least about 70, 80, 85, 90, 95, 96, 97, 98, or 99 percent identical to the amino acid sequence of SEQ ID NO: 2, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2 as determined using a computer program such as GAP, BLASTP, BLASTN, FASTA, BLASTA, BLASTX Bestfit, and the Smith-Waterman algorithm;

(c) a fragment of the amino acid sequence set forth in SEQ ID NO: 2 comprising at least about 25 amino acid residues, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(d) an amino acid sequence for an allelic variant or splice variant of either the amino acid sequence as set forth in SEQ ID NO: 2, or at least one of (a)-(b) wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2.

The invention further provides for an isolated polypeptide comprising the amino acid sequence selected from the group consisting of:

(a) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one conservative amino acid substitution, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(b) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one amino acid insertion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

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(c) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one amino acid deletion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

5 (d) the amino acid sequence as set forth in SEQ ID NO: 2 which has a C- and/or N-terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

10 (e) the amino acid sequence as set forth in SEQ ID NO: 2, with at least one modification selected from the group consisting of amino acid substitutions, amino acid insertions, amino acid deletions, C-terminal truncation, and N-terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2.

15 Also provided are fusion polypeptides comprising the amino acid sequences of (a)-(e) above.

The present invention also provides for an expression vector comprising the isolated nucleic acid molecules as set forth herein, recombinant host cells comprising recombinant  
20 nucleic acid molecules as set forth herein, and a method of producing an IL-17 receptor like polypeptide comprising culturing the host cells and optionally isolating the polypeptide so produced. These expression vectors include baculovirus expression vectors which utilize insect cells for  
25 expression.

A transgenic non-human animal comprising a nucleic acid molecule encoding an IL-17 receptor like polypeptide is also encompassed by the invention. The IL-17 receptor like nucleic acid molecules are introduced into the animal in a manner that  
30 allows expression and increased levels of the IL-17 receptor like polypeptide, which may include increased circulating levels. The transgenic non-human animal is preferably a mammal. Also provided is a transgenic non-human animals

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comprising a disruption in the nucleic acid molecule encoding a IL-17 receptor like polypeptide, which will knock-out or specifically decrease expression of the IL-17 receptor like polypeptide.

5 Also provided are derivatives of the IL-17 receptor like polypeptides of the present invention.

Analogs of the IL-17 receptor like polypeptides are provided for in the present invention which result from conservative and/or non-conservative amino acids substitutions  
10 of the IL-17 receptor like polypeptides of SEQ ID NO: 2. Such analogs include an IL-17 receptor like polypeptide wherein, for example the amino acid at position 45 of SEQ ID NO: 2 is glycine, proline or alanine, the amino acid at position 227 of SEQ ID NO: 2 is phenylalanine, leucine, valine, isoleucine,  
15 alanine or tyrosine, the amino acid at position 363 of SEQ ID NO: 2 is serine, threonine, alanine or cysteine, the amino acid at position 374 of SEQ ID NO: 2 is valine, isoleucine, leucine, phenylalanine, alanine or norleucine, the amino acid at position 515 of SEQ ID NO: 2 is aspartic acid or glutamic  
20 acid, the amino acid at position 6002 of SEQ ID NO: 2 is cysteine, serine or alanine.

Additionally provided are selective binding agents such as antibodies and peptides capable of specifically binding the IL-17 receptor like polypeptides of the invention. Such  
25 antibodies, polypeptides, peptides and small molecules may be agonistic or antagonistic.

Pharmaceutical compositions comprising the nucleotides, polypeptides, or selective binding agents of the present invention and one or more pharmaceutically acceptable  
30 formulation agents are also encompassed by the invention. The pharmaceutical compositions are used to provide therapeutically effective amounts of the nucleotides or polypeptides of the present invention. The invention is also

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directed to methods of using the polypeptides, nucleic acid molecules, and selective binding agents.

5 The IL-17 receptor like polypeptides and nucleic acid molecules of the present invention may be used to treat, prevent, ameliorate, diagnose and/or detect diseases and disorders, including those recited herein. Expression analysis in biological, cellular, or tissue samples suggests that IL-17 receptor like polypeptide may play a role in the diagnosis and/or treatment of the pathological conditions described herein. This expression can be detected with a  
10 diagnostic agent such as a IL-17 receptor like polynucleotide.

The invention encompasses diagnosing a pathological condition or the susceptibility to a pathological condition in a subject caused by or resulting from abnormal (*i.e.* increased  
15 or decreased) levels of IL-17 receptor like polypeptide comprising determining the presence or amount of expression of the IL-17 receptor like polypeptide in a sample and comparing the level of said polypeptide in a biological, tissue or cellular sample form either normal subjects at an earlier  
20 time, wherein susceptibility to a a pathological condition is based on the presence or amount of expression of the polypeptide.

The present invention also provides a method of assaying test molecules to identify a test molecule which binds to an  
25 IL-17 receptor like polypeptide. The method comprises contacting an IL-17 receptor like polypeptide with a test molecule and determining the extent of binding of the test molecule to the polypeptide. The method further comprises determining whether such test molecules are agonists or  
30 antagonists of an IL-17 receptor like polypeptide. The present invention further provides a method of testing the impact of molecules on the expression of IL-17 receptor like polypeptide or on the activity of IL-17 receptor like polypeptide.

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The present invention provides for methods of identifying antagonists and agonists of IL-17 receptor like biological activity comprising contacting a small molecule compound with IL-17 receptor like polypeptide and measuring IL-17 receptor like biological activity in the presence and absence of these small molecules. These small molecules can be a naturally occurring medicinal compound derived from combinational chemical libraries. In certain embodiments, an IL-17 receptor like polypeptide agonist or antagonist may be a protein, peptide, carbohydrate, lipid, or small molecule which interacts with a IL-17 receptor like polypeptide to regulate its activity.

Methods of regulating expression and modulating (i.e., increasing or decreasing) levels of an IL-17 receptor like polypeptide are also encompassed by the invention. One method comprises administering to an animal a nucleic acid molecule encoding an IL-17 receptor like polypeptide. In another method, a nucleic acid molecule comprising elements that regulate or modulate the expression of an IL-17 receptor like polypeptide may be administered. Examples of these methods include gene therapy, cell therapy, and anti-sense therapy as further described herein.

The IL-17 receptor like polypeptide can be used for identifying ligands thereof. Various forms of "expression cloning" have been used for cloning ligands for receptors. See e.g., Davis et al., *Cell*, 87:1161-1169 (1996). These and other IL-17 receptor like ligand cloning experiments are described in greater detail herein. Isolation of the IL-17 receptor like ligand(s) allows for the identification or development of novel agonists and/or antagonists of the IL-17 receptor like signaling pathway.

Agonists and antagonists include, but are not limited to, ligands to the IL-17 receptor like polypeptides, soluble IL-17

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receptor like polypeptides, anti-IL-17 receptor like selective binding agents (such as antibodies and derivatives thereof), small molecules, peptides and derivatives thereof, and antisense oligonucleotides, any of which can be used for  
5 treating one or more disease or disorder, including those disclosed herein.

The invention further encompasses methods for determining the presence of IL-17 receptor like nucleic acids in a biological, tissue or cellular sample. These methods comprise  
10 the steps of providing a biological sample suspected of containing IL-17 receptor like nucleic acids; contacting the sample with a diagnostic reagent of the present invention; detecting hybridization between nucleic acid in the biological sample and the diagnostic reagent; and comparing the level of  
15 hybridization between the biological sample with the level of hybridization between a known concentration of IL-17 receptor like nucleic acid and the diagnostic reagent. The polynucleotide detected in these methods may be an IL-17 receptor like DNA and/or a IL-17 receptor like RNA.

20 The invention also provides for a device which comprises a membrane for implantation in a patient; and cells encapsulated within said membrane, wherein said membrane is permeable to the protein product and impermeable to materials detrimental to said cells. The invention further provides for  
25 a device which comprises a membrane suitable for implantation and the IL-17 receptor like polypeptide encapsulated in a membrane that is permeable to the polypeptide.

#### Brief Description of the Figures

Figure 1 depicts a nucleic acid sequence (SEQ ID NO:1)  
30 and amino acid sequence (SEQ ID NO: 2) of the human IL-17 receptor like polypeptide.

Figure 2 depicts an overlap of the present human IL-17 receptor like (hIL-17RL) polypeptide and a known IL-17

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receptor family member (SEQ ID NO: 3).

### Detailed Description of the Invention

5 The section headings used herein are for organizational purposes only and are not to be construed as limiting the subject matter described therein. All references cited in this application are expressly incorporated by reference herein.

#### 10 Definitions

The terms "IL-17 receptor like gene" or "IL-17 receptor like nucleic acid molecule" or "polynucleotide" refers to a nucleic acid molecule comprising or consisting of a nucleotide sequence as set forth in SEQ ID NO: 1, a nucleotide sequence encoding the polypeptide as set forth in SEQ ID NO: 2, a  
15 nucleotide sequence of the DNA insert in Amgen deposit no. A-672-P, and nucleic acid molecules as defined herein.

The term "IL-17 receptor like polypeptide" refers to a polypeptide comprising the amino acid sequence of SEQ ID NO:  
20 2, and related polypeptides. Related polypeptides include: IL-17 receptor like polypeptide allelic variants, IL-17 receptor like polypeptide orthologs, IL-17 receptor like polypeptide splice variants, IL-17 receptor like polypeptide variants and IL-17 receptor like polypeptide derivatives. IL-  
25 17 receptor like polypeptides may be mature polypeptides, as defined herein, and may or may not have an amino terminal methionine residue, depending on the method by which they are prepared.

The term "IL-17 receptor like polypeptide allelic  
30 variant" refers to one of several possible naturally occurring alternate forms of a gene occupying a given locus on a chromosome of an organism or a population of organisms.



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The term "IL-17 receptor like polypeptide derivatives" refers to the polypeptide as set forth in SEQ ID NO: 2, IL-17 receptor like polypeptide allelic variants, IL-17 receptor like polypeptide orthologs, IL-17 receptor like polypeptide splice variants, or IL-17 receptor like polypeptide variants, as defined herein, that have been chemically modified.

The term "IL-17 receptor like polypeptide fragment" refers to a polypeptide that comprises a truncation at the amino terminus (with or without a leader sequence) and/or a truncation at the carboxy terminus of the polypeptide as set forth in SEQ ID NO: 2, IL-17 receptor like polypeptide allelic variants, IL-17 receptor like polypeptide orthologs, IL-17 receptor like polypeptide splice variants and/or an IL-17 receptor like polypeptide variant having one or more amino acid additions or substitutions or internal deletions (wherein the resulting polypeptide is at least 6 amino acids or more in length) as compared to the IL-17 receptor like polypeptide amino acid sequence set forth in SEQ ID NO: 2. IL-17 receptor like polypeptide fragments may result from alternative RNA splicing or from in vivo protease activity. For transmembrane or membrane-bound forms of an IL-17 receptor like polypeptide, preferred fragments include soluble forms such as those lacking a transmembrane or membrane-binding domain. In preferred embodiments, truncations comprise about 10 amino acids, or about 20 amino acids, or about 50 amino acids, or about 75 amino acids, or about 100 amino acids, or more than about 100 amino acids. The polypeptide fragments so produced will comprise about 25 contiguous amino acids, or about 50 amino acids, or about 75 amino acids, or about 100 amino acids, or about 150 amino acids, or about 200 amino acids. Such IL-17 receptor like polypeptide fragments may optionally comprise an amino terminal methionine residue. It will be appreciated that such fragments can be used, for example, to generate antibodies to IL-17 receptor like polypeptides.

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The term "IL-17 receptor like fusion polypeptide" refers to a fusion of one or more amino acids (such as a heterologous peptide or polypeptide) at the amino or carboxy terminus of the polypeptide as set forth in SEQ ID NO: 2, IL-17 receptor like polypeptide allelic variants, IL-17 receptor like polypeptide orthologs, IL-17 receptor like polypeptide splice variants, or IL-17 receptor like polypeptide variants having one or more amino acid deletions, substitutions or internal additions as compared to the IL-17 receptor like polypeptide amino acid sequence set forth in SEQ ID NO: 2.

The term "IL-17 receptor like polypeptide ortholog" refers to a polypeptide from another species that corresponds to IL-17 receptor like polypeptide amino acid sequence as set forth in SEQ ID NO: 2. For example, mouse and human IL-17 receptor like polypeptides are considered orthologs of each other.

The term "IL-17 receptor like polypeptide splice variant" refers to a nucleic acid molecule, usually RNA, which is generated by alternative processing of intron sequences in an RNA transcript of IL-17 receptor like polypeptide amino acid sequence as set forth in SEQ ID NO: 2.

The term "IL-17 receptor like polypeptide variants" refers to IL-17 receptor like polypeptides comprising amino acid sequences having one or more amino acid sequence substitutions, deletions (such as internal deletions and/or IL-17 receptor like polypeptide fragments), and/or additions (such as internal additions and/or IL-17 receptor like fusion polypeptides) as compared to the IL-17 receptor like polypeptide amino acid sequence set forth in SEQ ID NO: 2 (with or without a leader sequence). Variants may be naturally occurring (e.g., IL-17 receptor like polypeptide allelic variants, IL-17 receptor like polypeptide orthologs and IL-17 receptor like polypeptide splice variants) or

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artificially constructed. Such IL-17 receptor like polypeptide variants may be prepared from the corresponding nucleic acid molecules having a DNA sequence that varies accordingly from the DNA sequence as set forth in SEQ ID NO:

5 1. In preferred embodiments, the variants have from 1 to 3, or from 1 to 5, or from 1 to 10, or from 1 to 15, or from 1 to 20, or from 1 to 25, or from 1 to 50, or from 1 to 75, or from 1 to 100, or more than 100 amino acid substitutions, insertions, additions and/or deletions, wherein the  
10 substitutions may be conservative, or non-conservative, or any combination thereof.

The term "antigen" refers to a molecule or a portion of a molecule capable of being bound by a selective binding agent, such as an antibody, and additionally capable of being  
15 used in an animal to produce antibodies capable of binding to an epitope of that antigen. An antigen may have one or more epitopes.

The term specific binding reaction referred to above is meant to indicate that the antigen will react, in a highly  
20 selective manner, with its corresponding antibody and not with the multitude of other antibodies which can be evoked by other antigens.

The term "biologically active IL-17 receptor like polypeptides" refers to IL-17 receptor like polypeptides,  
25 including fragments, variants, derivatives, having at least one activity characteristic of the polypeptide comprising the amino acid sequence of SEQ ID NO: 2. In addition, a IL-17 receptor like polypeptide may be active as an immunogen, that is, the polypeptide contains at least one epitope to which  
30 antibodies may be raised.

The terms "effective amount" and "therapeutically effective amount" each refer to the amount of a IL-17 receptor like polypeptide or IL-17 receptor like nucleic acid molecule

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used to support an observable level of one or more biological activities of the IL-17 receptor like polypeptides as set forth herein.

5 The term "expression vector" refers to a vector which is suitable for use in a host cell and contains nucleic acid sequences which direct and/or control the expression of inserted heterologous nucleic acid sequences. Expression includes, but is not limited to, processes such as transcription, translation, and RNA splicing, if introns are  
10 present.

The term "host cell" is used to refer to a cell which has been transformed, or is capable of being transformed with a nucleic acid sequence and then of expressing a selected gene of interest. The term includes the progeny of the parent  
15 cell, whether or not the progeny is identical in morphology or in genetic make-up to the original parent, so long as the selected gene is present.

The term "identity" as known in the art, refers to a relationship between the sequences of two or more polypeptide  
20 molecules or two or more nucleic acid molecules, as determined by comparing the sequences. In the art, "identity" also means the degree of sequence relatedness between nucleic acid molecules or polypeptides, as the case may be, as determined by the match between strings of two or more nucleotide or two  
25 or more amino acid sequences. "Identity" measures the percent of identical matches between the smaller of two or more sequences with gap alignments (if any) addressed by a particular mathematical model or computer program (i.e., "algorithms").

30 The term "similarity" is a related concept, but in contrast to "identity", refers to a measure of similarity which includes both identical matches and conservative substitution matches. If two polypeptide sequences have, for

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example, 10/20 identical amino acids, and the remainder are all non-conservative substitutions, then the percent identity and similarity would both be 50%. If in the same example, there are 5 more positions where there are conservative substitutions, then the percent identity remains 50%, but the per cent similarity would be 75% (15/20). Therefore, in cases where there are conservative substitutions, the degree of similarity between two polypeptides will be higher than the percent identity between those two polypeptides.

10 The term "isolated nucleic acid molecule" refers to a nucleic acid molecule of the invention that (1) has been separated from at least about 50 percent of proteins, lipids, carbohydrates or other materials with which it is naturally found when total DNA is isolated from the source cells, (2) is  
15 not linked to all or a portion of a polynucleotide to which the "isolated nucleic acid molecule" is linked in nature, (3) is operably linked to a polynucleotide which it is not linked to in nature, or (4) does not occur in nature as part of a larger polynucleotide sequence. Preferably, the isolated  
20 nucleic acid molecule of the present invention is substantially free from at least one contaminating nucleic acid molecule with which it is naturally associated. Preferably, the isolated nucleic acid molecule of the present invention is substantially free from any other contaminating  
25 nucleic acid molecule(s) or other contaminants that are found in its natural environment that would interfere with its use in polypeptide production or its therapeutic, diagnostic, prophylactic or research use.

The term "isolated polypeptide" refers to a polypeptide  
30 of the present invention that (1) has been separated from at least about 50 percent of polynucleotides, lipids, carbohydrates or other materials with which it is naturally found when isolated from the source cell, (2) is not linked (by covalent or noncovalent interaction) to all or a portion

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of a polypeptide to which the "isolated polypeptide" is linked to in nature, (3) is operably linked (by covalent or noncovalent interaction) to a polypeptide with which it is not linked in nature, or (4) does not occur in nature.

5 Preferably, is free from at least one contaminating polypeptide or other contaminants that are found in its natural environment. Preferably, the isolated polypeptide is substantially free from any other contaminating polypeptides or other contaminants that are found in its natural

10 environment which would interfere with its therapeutic, diagnostic, prophylactic or research use.

The term "mature IL-17 receptor like polypeptide" refers to an IL-17 receptor like polypeptide lacking a leader sequence. A mature IL-17 receptor like polypeptide may also

15 include other modifications such as proteolytic processing of the amino terminus (with or without a leader sequence) and/or the carboxy terminus, cleavage of a smaller polypeptide from a larger precursor, N-linked and/or O-linked glycosylation, and the like.

20 The term "nucleic acid sequence" or "nucleic acid molecule" refers to a DNA or RNA sequence. The term encompasses molecules formed from any of the known base analogs of DNA and RNA such as, but not limited to 4-acetylcytosine, 8-hydroxy-N6-methyladenosine, aziridinyl-

25 cytosine, pseudoisocytosine, 5-(carboxyhydroxymethyl) uracil, 5-fluorouracil, 5-bromouracil, 5-carboxymethylaminomethyl-2-thiouracil, 5-carboxy-methylaminomethyluracil, dihydrouracil, inosine, N6-iso-pentenyladenine, 1-methyladenine, 1-methylpseudouracil, 1-methylguanine, 1-methylinosine, 2,2-

30 dimethyl-guanine, 2-methyladenine, 2-methylguanine, 3-methylcytosine, 5-methylcytosine, N6-methyladenine, 7-methylguanine, 5-methylaminomethyluracil, 5-methoxyamino-methyl-2-thiouracil, beta-D-mannosylqueosine, 5'-methoxycarbonyl-methyluracil, 5-methoxyuracil, 2-methylthio-

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N6-isopentenyladenine, uracil-5-oxyacetic acid methylester, uracil-5-oxyacetic acid, oxybutoxosine, pseudouracil, queosine, 2-thiocytosine, 5-methyl-2-thiouracil, 2-thiouracil, 4-thiouracil, 5-methyluracil, N-uracil-5-oxyacetic acid methylester, uracil-5-oxyacetic acid, pseudouracil, queosine, 2-thiocytosine, and 2,6-diaminopurine.

The term "naturally occurring" or "native" when used in connection with biological materials such as nucleic acid molecules, polypeptides, host cells, and the like, refers to materials which are found in nature and are not manipulated by man. Similarly, "non-naturally occurring" or "non-native" as used herein refers to a material that is not found in nature or that has been structurally modified or synthesized by man.

The term "operably linked" is used herein to refer to an arrangement of flanking sequences wherein the flanking sequences so described are configured or assembled so as to perform their usual function. Thus, a flanking sequence operably linked to a coding sequence may be capable of effecting the replication, transcription and/or translation of the coding sequence. For example, a coding sequence is operably linked to a promoter when the promoter is capable of directing transcription of that coding sequence. A flanking sequence need not be contiguous with the coding sequence, so long as it functions correctly. Thus, for example, intervening untranslated yet transcribed sequences can be present between a promoter sequence and the coding sequence and the promoter sequence can still be considered "operably linked" to the coding sequence.

The term "pharmaceutically acceptable carrier" or "physiologically acceptable carrier" as used herein refers to one or more formulation materials suitable for accomplishing or enhancing the delivery of the IL-17 receptor like polypeptide, IL-17 receptor like nucleic acid molecule or IL-

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17 receptor like selective binding agent as a pharmaceutical composition.

The term "selective binding agent" refers to a molecule or molecules having specificity for an IL-17 receptor like polypeptide. Selective binding agents include antibodies, such as polyclonal antibodies, monoclonal antibodies (mAbs), chimeric antibodies, CDR-grafted antibodies, anti-idiotypic (anti-Id) antibodies to antibodies that can be labeled in soluble or bound forms, as well as fragments, regions, or derivatives thereof which are provided by known techniques, including, but not limited to enzymatic cleavage, peptide synthesis, or recombinant techniques. The anti-IL-17 receptor like selective binding agents of the present invention are capable, for example, of binding portions of IL-17 like molecules to IL-17 receptor like polypeptides of the present invention.

As used herein, the terms, "specific" and "specificity" refer to the ability of the selective binding agents to bind to human IL-17 receptor like polypeptides and not to bind to human non-IL-17 receptor like polypeptides. It will be appreciated, however, that the selective binding agents may also bind orthologs of the polypeptide as set forth in SEQ ID NO: 2, that is, interspecies versions thereof, such as mouse and rat polypeptides.

The term "transduction" is used to refer to the transfer of genes from one bacterium to another, usually by a phage. "Transduction" also refers to the acquisition and transfer of eukaryotic cellular sequences by retroviruses.

The term "transfection" is used to refer to the uptake of foreign or exogenous DNA by a cell, and a cell has been "transfected" when the exogenous DNA has been introduced inside the cell membrane. A number of transfection techniques are well known in the art and are disclosed herein. See, for



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example, Graham et al., *Virology*, 52:456 (1973); Sambrook et al., *Molecular Cloning, a laboratory Manual*, Cold Spring Harbor Laboratories (New York, 1989); Davis et al., *Basic Methods in Molecular Biology*, Elsevier, 1986; and Chu et al., *Gene*, 13:197 (1981). Such techniques can be used to introduce one or more exogenous DNA moieties into suitable host cells.

The term "transformation" as used herein refers to a change in a cell's genetic characteristics, and a cell has been transformed when it has been modified to contain a new DNA. For example, a cell is transformed where it is genetically modified from its native state. Following transfection or transduction, the transforming DNA may recombine with that of the cell by physically integrating into a chromosome of the cell, may be maintained transiently as an episomal element without being replicated, or may replicate independently as a plasmid. A cell is considered to have been stably transformed when the DNA is replicated with the division of the cell.

The term "vector" is used to refer to any molecule (e.g., nucleic acid, plasmid, or virus) used to transfer coding information to a host cell.

#### Relatedness of Nucleic Acid Molecules and/or Polypeptides

It is understood that related nucleic acid molecules include allelic or splice variants of the nucleic acid molecule of SEQ ID NO: 1, and include sequences which are complementary to any of the above nucleotide sequences. Related nucleic acid molecules also include a nucleotide sequence encoding a polypeptide comprising or consisting essentially of a substitution, modification, addition and/or a deletion of one or more amino acid residues compared to the polypeptide in SEQ ID NO: 2.

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Fragments include molecules which encode a polypeptide of at least about 25 amino acid residues, or about 50, or about 75, or about 100, or greater than about 100 amino acid residues of the polypeptide of SEQ ID NO: 2.

5 In addition, related IL-17 receptor like nucleic acid molecules include those molecules which comprise nucleotide sequences which hybridize under moderately or highly stringent conditions as defined herein with the fully complementary sequence of the nucleic acid molecule of SEQ ID NO: 1, or of a  
10 molecule encoding a polypeptide, which polypeptide comprises the amino acid sequence as shown in SEQ ID NO: 2, or of a nucleic acid fragment as defined herein, or of a nucleic acid fragment encoding a polypeptide as defined herein. Hybridization probes may be prepared using the IL-17 receptor  
15 like sequences provided herein to screen cDNA, genomic or synthetic DNA libraries for related sequences. Regions of the DNA and/or amino acid sequence of IL-17 receptor like polypeptide that exhibit significant identity to known sequences are readily determined using sequence alignment  
20 algorithms as described herein and those regions may be used to design probes for screening.

The term "highly stringent conditions" refers to those conditions that are designed to permit hybridization of DNA strands whose sequences are highly complementary, and to  
25 exclude hybridization of significantly mismatched DNAs. Hybridization stringency is principally determined by temperature, ionic strength, and the concentration of denaturing agents such as formamide. Examples of "highly stringent conditions" for hybridization and washing are 0.015M  
30 sodium chloride, 0.0015M sodium citrate at 65-68°C or 0.015M sodium chloride, 0.0015M sodium citrate, and 50% formamide at 42°C. See Sambrook, Fritsch & Maniatis, Molecular Cloning: A Laboratory Manual, 2<sup>nd</sup> Ed., Cold Spring Harbor Laboratory, (Cold Spring Harbor, N.Y. 1989); Anderson et al., Nucleic Acid

Hybridisation: a practical approach, Ch. 4, IRL Press Limited (Oxford, England).

More stringent conditions (such as higher temperature, lower ionic strength, higher formamide, or other denaturing agent) may also be used, however, the rate of hybridization will be affected. Other agents may be included in the hybridization and washing buffers for the purpose of reducing non-specific and/or background hybridization. Examples are 0.1% bovine serum albumin, 0.1% polyvinyl-pyrrolidone, 0.1% sodium pyrophosphate, 0.1% sodium dodecylsulfate (NaDodSO<sub>4</sub> or SDS), ficoll, Denhardt's solution, sonicated salmon sperm DNA (or other non-complementary DNA), and dextran sulfate, although other suitable agents can also be used. The concentration and types of these additives can be changed without substantially affecting the stringency of the hybridization conditions. Hybridization experiments are usually carried out at pH 6.8-7.4, however, at typical ionic strength conditions, the rate of hybridization is nearly independent of pH. See Anderson et al., Nucleic Acid Hybridisation: a Practical Approach, Ch. 4, IRL Press Limited (Oxford, England).

Factors affecting the stability of a DNA duplex include base composition, length, and degree of base pair mismatch. Hybridization conditions can be adjusted by one skilled in the art in order to accommodate these variables and allow DNAs of different sequence relatedness to form hybrids. The melting temperature of a perfectly matched DNA duplex can be estimated by the following equation:

$$T_m(^{\circ}\text{C}) = 81.5 + 16.6(\log[\text{Na}^+]) + 0.41(\% \text{G+C}) - 600/N - 0.72(\% \text{formamide})$$

where N is the length of the duplex formed, [Na<sup>+</sup>] is the molar concentration of the sodium ion in the hybridization or washing solution, %G+C is the percentage of (guanine+cytosine) bases in the hybrid. For imperfectly matched hybrids, the

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melting temperature is reduced by approximately 1°C for each 1% mismatch.

The term "moderately stringent conditions" refers to conditions under which a DNA duplex with a greater degree of base pair mismatching than could occur under "highly stringent conditions" is able to form. Examples of typical "moderately stringent conditions" are 0.015M sodium chloride, 0.0015M sodium citrate at 50-65°C or 0.015M sodium chloride, 0.0015M sodium citrate, and 20% formamide at 37-50°C. By way of example, a "moderately stringent" condition of 50°C in 0.015 M sodium ion will allow about a 21% mismatch.

It will be appreciated by those skilled in the art that there is no absolute distinction between "highly" and "moderately" stringent conditions. For example, at 0.015M sodium ion (no formamide), the melting temperature of perfectly matched long DNA is about 71°C. With a wash at 65°C (at the same ionic strength), this would allow for approximately a 6% mismatch. To capture more distantly related sequences, one skilled in the art can simply lower the temperature or raise the ionic strength.

A good estimate of the melting temperature in 1M NaCl\* for oligonucleotide probes up to about 20nt is given by:

$$T_m = 2^\circ\text{C per A-T base pair} + 4^\circ\text{C per G-C base pair}$$

\*The sodium ion concentration in 6X salt sodium citrate (SSC) is 1M. See Suggs et al., Developmental Biology Using Purified Genes, p. 683, Brown and Fox (eds.) (1981).

High stringency washing conditions for oligonucleotides are usually at a temperature of 0-5°C below the  $T_m$  of the oligonucleotide in 6X SSC, 0.1% SDS.

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In another embodiment, related nucleic acid molecules comprise or consist of a nucleotide sequence that is about 70 percent identical to the nucleotide sequence as shown in SEQ ID NO: 1, or comprise or consist essentially of a nucleotide sequence encoding a polypeptide that is about 70 percent identical to the polypeptide as set forth in SEQ ID NO: 2. In preferred embodiments, the nucleotide sequences are about 75 percent, or about 80 percent, or about 85 percent, or about 90 percent, or about 95, 96, 97, 98, or 99 percent identical to the nucleotide sequence as shown in SEQ ID NO: 1, or the nucleotide sequences encode a polypeptide that is about 75 percent, or about 80 percent, or about 85 percent, or about 90 percent, or about 95, 96, 97, 98, or 99 percent identical to the polypeptide sequence as set forth in SEQ ID NO: 2.

15 Differences in the nucleic acid sequence may result in conservative and/or non-conservative modifications of the amino acid sequence relative to the amino acid sequence of SEQ ID NO: 2.

Conservative modifications to the amino acid sequence of SEQ ID NO: 2 (and the corresponding modifications to the encoding nucleotides) will produce IL-17 receptor like polypeptides having functional and chemical characteristics similar to those of naturally occurring IL-17 receptor like polypeptide. In contrast, substantial modifications in the functional and/or chemical characteristics of IL-17 receptor like polypeptides may be accomplished by selecting substitutions in the amino acid sequence of SEQ ID NO: 2 that differ significantly in their effect on maintaining (a) the structure of the molecular backbone in the area of the substitution, for example, as a sheet or helical conformation, (b) the charge or hydrophobicity of the molecule at the target site, or (c) the bulk of the side chain.

For example, a "conservative amino acid substitution" may involve a substitution of a native amino acid residue with a

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nonnative residue such that there is little or no effect on the polarity or charge of the amino acid residue at that position. Furthermore, any native residue in the polypeptide may also be substituted with alanine, as has been previously  
5 described for "alanine scanning mutagenesis."

Conservative amino acid substitutions also encompass non-naturally occurring amino acid residues which are typically incorporated by chemical peptide synthesis rather than by  
10 synthesis in biological systems. These include peptidomimetics, and other reversed or inverted forms of amino acid moieties. It will be appreciated by those of skill in the art that nucleic acid polypeptide molecules described herein may be chemically synthesized as well as produced by  
15 recombinant means.

Naturally occurring residues may be divided into classes based on common side chain properties:

- 1) hydrophobic: norleucine, Met, Ala, Val, Leu, Ile;
- 2) neutral hydrophilic: Cys, Ser, Thr, Asn, Gln;
- 20 3) acidic: Asp, Glu;
- 4) basic: His, Lys, Arg;
- 5) residues that influence chain orientation: Gly, Pro; and
- 6) aromatic: Trp, Tyr, Phe.

25 For example, non-conservative substitutions may involve the exchange of a member of one of these classes for a member from another class. Such substituted residues may be introduced into regions of the human IL-17 receptor like polypeptide that are homologous with non-human IL-17 receptor  
30 like polypeptide orthologs, or into the non-homologous regions of the molecule.

In making such changes, the hydropathic index of amino acids may be considered. Each amino acid has been assigned a

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hydropathic index on the basis of their hydrophobicity and charge characteristics, these are: isoleucine (+4.5); valine (+4.2); leucine (+3.8); phenylalanine (+2.8); cysteine/cystine (+2.5); methionine (+1.9); alanine (+1.8); glycine (-0.4);  
5 threonine (-0.7); serine (-0.8); tryptophan (-0.9); tyrosine (-1.3); proline (-1.6); histidine (-3.2); glutamate (-3.5); glutamine (-3.5); aspartate (-3.5); asparagine (-3.5); lysine (-3.9); and arginine (-4.5).

The importance of the hydropathic amino acid index in  
10 conferring interactive biological function on a protein is understood in the art. Kyte et al., *J. Mol. Biol.*, 157:105-131 (1982). It is known that certain amino acids may be substituted for other amino acids having a similar hydropathic index or score and still retain a similar biological activity.  
15 In making changes based upon the hydropathic index, the substitution of amino acids whose hydropathic indices are within  $\pm 2$  is preferred, those which are within  $\pm 1$  are particularly preferred, and those within  $\pm 0.5$  are even more particularly preferred.

20 It is also understood in the art that the substitution of like amino acids can be made effectively on the basis of hydrophilicity, particularly where the biologically functionally equivalent protein or peptide thereby created is intended for use in immunological embodiments, as in the  
25 present case. The greatest local average hydrophilicity of a protein, as governed by the hydrophilicity of its adjacent amino acids, correlates with its immunogenicity and antigenicity, i.e., with a biological property of the protein.

The following hydrophilicity values have been assigned to  
30 amino acid residues: arginine (+3.0); lysine (+3.0); aspartate (+3.0  $\pm$  1); glutamate (+3.0  $\pm$  1); serine (+0.3); asparagine (+0.2); glutamine (+0.2); glycine (0); threonine (-0.4); proline (-0.5  $\pm$  1); alanine (-0.5); histidine (-0.5); cysteine

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(-1.0); methionine (-1.3); valine (-1.5); leucine (-1.8); isoleucine (-1.8); tyrosine (-2.3); phenylalanine (-2.5); tryptophan (-3.4). In making changes based upon similar hydrophilicity values, the substitution of amino acids whose hydrophilicity values are within  $\pm 2$  is preferred, those which are within  $\pm 1$  are particularly preferred, and those within  $\pm 0.5$  are even more particularly preferred. One may also identify epitopes from primary amino acid sequences on the basis of hydrophilicity. These regions are also referred to as "epitopic core regions."

Desired amino acid substitutions (whether conservative or non-conservative) can be determined by those skilled in the art at the time such substitutions are desired. For example, amino acid substitutions can be used to identify important residues of the IL-17 receptor like polypeptide, or to increase or decrease the affinity of the IL-17 receptor like polypeptides described herein.

Exemplary amino acid substitutions are set forth in Table

I.



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Table IAmino Acid Substitutions

Original Residues	Exemplary Substitutions	Preferred Substitutions
Ala	Val, Leu, Ile	Val
Arg	Lys, Gln, Asn	Lys
Asn	Gln	Gln
Asp	Glu	Glu
Cys	Ser, Ala	Ser
Gln	Asn	Asn
Glu	Asp	Asp
Gly	Pro, Ala	Ala
His	Asn, Gln, Lys, Arg	Arg
Ile	Leu, Val, Met, Ala, Phe, Norleucine	Leu
Leu	Norleucine, Ile, Val, Met, Ala, Phe	Ile
Lys	Arg, 1,4 Diamino-butyrac Acid, Gln, Asn	Arg
Met	Leu, Phe, Ile	Leu
Phe	Leu, Val, Ile, Ala, Tyr	Leu
Pro	Ala	Gly
Ser	Thr, Ala, Cys	Thr
Thr	Ser	Ser
Trp	Tyr, Phe	Tyr
Tyr	Trp, Phe, Thr, Ser	Phe
Val	Ile, Met, Leu, Phe, Ala, Norleucine	Leu

5 A skilled artisan will be able to determine suitable variants of the polypeptide as set forth in SEQ ID NO: 2 using well known techniques. For identifying suitable areas of the

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molecule that may be changed without destroying biological activity. Also, one skilled in the art will realize that even areas that may be important for biological activity or for structure may be subject to conservative amino acid  
5 substitutions without destroying the biological activity or without adversely affecting the polypeptide structure.

For example, when similar polypeptides with similar activities from the same species or from other species are known, one skilled in the art may compare the amino acid  
10 sequence of an IL-17 receptor like polypeptide to such similar polypeptides. With such a comparison, one can identify residues and portions of the molecules that are conserved among similar polypeptides. It will be appreciated that  
15 changes in areas of an IL-17 receptor like polypeptide that are not conserved relative to such similar polypeptides would be less likely to adversely affect the biological activity and/or structure of the IL-17 receptor like polypeptide. One  
20 skilled in the art would also know that, even in relatively conserved regions, one may substitute chemically similar amino acids for the naturally occurring residues while retaining activity (conservative amino acid residue substitutions).  
Therefore, even areas that may be important for biological activity or for structure may be subject to conservative amino acid substitutions without destroying the biological activity  
25 or without adversely affecting the polypeptide structure.

For predicting suitable areas of the molecule that may be changed without destroying activity, one skilled in the art may target areas not believed to be important for activity. For example, when similar polypeptides with similar activities  
30 from the same species or from other species are known, one skilled in the art may compare the amino acid sequence of IL-17 receptor like polypeptide to such similar polypeptides. After making such a comparison, one skilled in the art can

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determine residues and portions of the molecules that are conserved among similar polypeptides. One skilled in the art would know that changes in areas of the IL-17 receptor like molecule that are not conserved would be less likely to adversely affect the biological activity and/or structure of a IL-17 receptor like polypeptide. One skilled in the art would also know that, even in relatively conserved regions, one may substitute chemically similar amino acids for the naturally occurring residues while retaining activity (conservative amino acid residue substitutions).

Additionally, one skilled in the art can review structure-function studies identifying residues in similar polypeptides that are important for activity or structure. In view of such a comparison, one can predict the importance of amino acid residues in an IL-17 receptor like polypeptide that correspond to amino acid residues that are important for activity or structure in similar polypeptides. One skilled in the art may opt for chemically similar amino acid substitutions for such predicted important amino acid residues of IL-17 receptor like polypeptides.

One skilled in the art can also analyze the three-dimensional structure and amino acid sequence in relation to that structure in similar polypeptides. In view of that information, one skilled in the art may predict the alignment of amino acid residues of an IL-17 receptor like polypeptide with respect to its three dimensional structure. One skilled in the art may choose not to make radical changes to amino acid residues predicted to be on the surface of the protein, since such residues may be involved in important interactions with other molecules. Moreover, one skilled in the art may generate test variants containing a single amino acid substitution at each desired amino acid residue. The variants can then be screened using activity assays known to those skilled in the art. Such variants could be used to gather

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information about suitable variants. For example, if one discovered that a change to a particular amino acid residue resulted in destroyed, undesirably reduced, or unsuitable activity, variants with such a change would be avoided. In other words, based on information gathered from such routine experiments, one skilled in the art can readily determine the amino acids where further substitutions should be avoided either alone or in combination with other mutations.

A number of scientific publications have been devoted to the prediction of secondary structure. See Moulton J., *Curr. Op. in Biotech.*, 7(4):422-427 (1996), Chou et al., *Biochemistry*, 13(2):222-245 (1974); Chou et al., *Biochemistry*, 113(2):211-222 (1974); Chou et al., *Adv. Enzymol. Relat. Areas Mol. Biol.*, 47:45-148 (1978); Chou et al., *Ann. Rev. Biochem.*, 47:251-276 and Chou et al., *Biophys. J.*, 26:367-384 (1979). Moreover, computer programs are currently available to assist with predicting secondary structure. One method of predicting secondary structure is based upon homology modeling. For example, two polypeptides or proteins which have a sequence identity of greater than 30%, or similarity greater than 40% often have similar structural topologies. The recent growth of the protein structural data base (PDB) has provided enhanced predictability of secondary structure, including the potential number of folds within a polypeptide's or protein's structure. See Holm et al., *Nucl. Acid. Res.*, 27(1):244-247 (1999). It has been suggested (Brenner et al., *Curr. Op. Struct. Biol.*, 7(3):369-376 (1997)) that there are a limited number of folds in a given polypeptide or protein and that once a critical number of structures have been resolved, structural prediction will gain dramatically in accuracy.

Additional methods of predicting secondary structure include "threading" (Jones, D., *Curr. Opin. Struct. Biol.*, 7(3):377-87 (1997); Sippl et al., *Structure*, 4(1):15-9 (1996)), "profile analysis" (Bowie et al., *Science*, 253:164-

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170 (1991); Gribskov et al., *Meth. Enzym.*, 183:146-159 (1990); Gribskov et al., *Proc. Nat. Acad. Sci.*, 84(13):4355-4358 (1987)), and "evolutionary linkage" (See Home, *supra*, and Brenner, *supra*).

5 IL-17 receptor like polypeptide analogs of the invention can be determined by comparing the amino acid sequence of IL-17 receptor like polypeptide with related family members. An exemplary IL-17 receptor like polypeptide related family member is a human IL-17 receptor. This comparison can be  
10 accomplished by using a Pileup alignment (Wisconsin GCG Program Package) or an equivalent (overlapping) comparison with multiple family members within conserved and non-conserved regions.

As shown in Figure 2, the predicted amino acid sequences  
15 of human IL-17 receptor like polypeptides (SEQ ID NO: 2) is aligned with a known human IL-17 receptor family member (SEQ ID NO: 3). Other IL-17 receptor like polypeptide analogs can be determined using these or other methods known to those of skill in the art. These overlapping sequences provide  
20 guidance for conservative and non-conservative amino acids substitutions resulting in additional IL-17 receptor like analogs. It will be appreciated that these amino acid substitutions can consist of naturally occurring or non-naturally occurring amino acids. For example, potential IL-17  
25 receptor like analogs may have the Gly at residue at position 45 of SEQ ID NO: 2 substituted with a Pro or Ala residue, the Phe residue at position 227 of SEQ ID NO: 2 substituted with a Leu, Val, Ile, Ala or Tyr residue, and/or the Ser residue at position 363 of SEQ ID NO: 2 substituted with a Thr, Ala or  
30 Cys. In addition, potential IL-17 receptor like analogs may have the Val residue at position 374 of SEQ ID NO: 2 substituted with a Ile, Met, Leu, Phe, Ala or norleucine residue, the Cys residue at position 385 of SEQ ID NO: 2, substituted with a Ser or Ala residue, the Asp residue at  
35 position 515 of SEQ ID NO: 2 substituted with a Glu residue

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and/or the Cys residue at position 602 substituted with a Ser or Ala.

Preferred IL-17 receptor like polypeptide variants include glycosylation variants wherein the number and/or type of glycosylation sites has been altered compared to the amino acid sequence set forth in SEQ ID NO: 2. In one embodiment, IL-17 receptor like polypeptide variants comprise a greater or a lesser number of N-linked glycosylation sites than the amino acid sequence set forth in SEQ ID NO: 2. An N-linked glycosylation site is characterized by the sequence: Asn-X-Ser or Asn-X-Thr, wherein the amino acid residue designated as X may be any amino acid residue except proline. The substitution(s) of amino acid residues to create this sequence provides a potential new site for the addition of an N-linked carbohydrate chain. Alternatively, substitutions which eliminate this sequence will remove an existing N-linked carbohydrate chain. Also provided is a rearrangement of N-linked carbohydrate chains wherein one or more N-linked glycosylation sites (typically those that are naturally occurring) are eliminated and one or more new N-linked sites are created. Additional preferred IL-17 receptor like variants include cysteine variants, wherein one or more cysteine residues are deleted from or substituted for another amino acid (e.g., serine) as compared to the amino acid sequence set forth in SEQ ID NO: 2. Cysteine variants are useful when IL-17 receptor like polypeptides must be refolded into a biologically active conformation such as after the isolation of insoluble inclusion bodies. Cysteine variants generally have fewer cysteine residues than the native protein, and typically have an even number to minimize interactions resulting from unpaired cysteines.

In addition, the polypeptide comprising the amino acid sequence of SEQ ID NO: 2 or an IL-17 receptor like polypeptide variant may be fused to a homologous polypeptide to form a

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homodimer or to a heterologous polypeptide to form a heterodimer. Heterologous peptides and polypeptides include, but are not limited to: an epitope to allow for the detection and/or isolation of an IL-17 receptor like fusion polypeptide; 5 a transmembrane receptor protein or a portion thereof, such as an extracellular domain, or a transmembrane and intracellular domain; a ligand or a portion thereof which binds to a transmembrane receptor protein; an enzyme or portion thereof which is catalytically active; a polypeptide or peptide which 10 promotes oligomerization, such as a leucine zipper domain; a polypeptide or peptide which increases stability, such as an immunoglobulin constant region; and a polypeptide which has a therapeutic activity different from the polypeptide comprising the amino acid sequence as set forth in SEQ ID NO: 2 or an IL- 15 17 receptor like polypeptide variant. In addition, a IL-17 receptor like polypeptide may be fused to itself or to a fragment, variant, or derivative thereof.

Fusions can be made either at the amino terminus or at the carboxy terminus of the polypeptide comprising the amino 20 acid sequence set forth in SEQ ID NO: 2 or an IL-17 receptor like polypeptide variant. Fusions may be direct with no linker or adapter molecule or indirect using a linker or adapter molecule. A linker or adapter molecule may be one or more amino acid residues, typically up to about 20 amino acids 25 residues, or up to about 50 amino acid residues. A linker or adapter molecule may also be designed with a cleavage site for a DNA restriction endonuclease or for a protease to allow for the separation of the fused moieties. It will be appreciated that once constructed, the fusion polypeptides can be 30 derivatized according to the methods described herein.

In a further embodiment of the invention, the polypeptide comprising the amino acid sequence of SEQ ID NO: 2 or an IL-17 receptor like polypeptide variant, including a fragment, variant and/or derivative, is fused to one or more domains of

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an Fc region of human IgG. Antibodies comprise two functionally independent parts, a variable domain known as "Fab", which binds antigen, and a constant domain known as "Fc", which is involved in effector functions such as complement activation and attack by phagocytic cells. An Fc has a long serum half-life, whereas an Fab is short-lived. Capon et al., *Nature*, 337:525-31 (1989). When constructed together with a therapeutic protein, an Fc domain can provide longer half-life or incorporate such functions as Fc receptor binding, protein A binding, complement fixation and perhaps even placental transfer. *Id.* Table II summarizes the use of certain Fc fusions known in the art, including materials and methods applicable to the production of fused IL-17 receptor like polypeptide.

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Table II  
Fc Fusion with Therapeutic Proteins

Form of Fc	Fusion partner	Therapeutic implications	Reference
IgG1	N-terminus of CD30-L	Hodgkin's disease; anaplastic lymphoma; T-cell leukemia	U.S. Patent No. 5,480,981
Murine Fcγ2a	IL-10	anti-inflammatory; transplant rejection	Zheng et al. (1995), <i>J. Immunol.</i> , <u>154</u> : 5590-5600
IgG1	TNF receptor	septic shock	Fisher et al. (1996), <i>N. Engl. J. Med.</i> , <u>334</u> : 1697-1702; Van Zee et al., (1996), <i>J. Immunol.</i> , <u>156</u> : 2221-2230
IgG, IgA, IgM, or IgE (excluding the first domain)	TNF receptor	inflammation, autoimmune disorders	U.S. Pat. No. 5,808,029, issued September 15, 1998
IgG1	CD4 receptor	AIDS	Capon et al. (1989), <i>Nature</i> <u>337</u> : 525-531
IgG1, IgG3	N-terminus of IL-2	anti-cancer, antiviral	Harvill et al. (1995), <i>Immunotech.</i> , <u>1</u> : 95-105
IgG1	C-terminus of OPG	osteoarthritis; bone density	WO 97/23614, published July 3, 1997
IgG1	N-terminus of leptin	anti-obesity	PCT/US 97/23183, filed December 11, 1997
Human Ig Cγ1	CTLA-4	autoimmune disorders	Linsley (1991), <i>J. Exp. Med.</i> , <u>174</u> : 561-569

In one example, all or a portion of the human IgG hinge, CH2 and CH3 regions may be fused at either the N-terminus or C-terminus of the IL-17 receptor like polypeptides using

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methods known to the skilled artisan. In another example, a portion of hinge regions and CH2 and CH3 regions may be fused. The resulting IL-17 receptor like fusion polypeptide may be purified by use of a Protein A affinity column. Peptides and proteins fused to an Fc region have been found to exhibit a substantially greater half-life in vivo than the unfused counterpart. Also, a fusion to an Fc region allows for dimerization/multimerization of the fusion polypeptide. The Fc region may be a naturally occurring Fc region, or may be altered to improve certain qualities, such as therapeutic qualities, circulation time, reduce aggregation, etc.

Identity and similarity of related nucleic acid molecules and polypeptides can be readily calculated by known methods. Such methods include, but are not limited to, those described in Computational Molecular Biology, Lesk, A.M., ed., Oxford University Press, New York, 1988; Biocomputing: Informatics and Genome Projects, Smith, D.W., ed., Academic Press, New York, 1993; Computer Analysis of Sequence Data, Part 1, Griffin, A.M., and Griffin, H.G., eds., Humana Press, New Jersey, 1994; Sequence Analysis in Molecular Biology, von Heinje, G., Academic Press, 1987; Sequence Analysis Primer, Gribskov, M. and Devereux, J., eds., M. Stockton Press, New York, 1991; and Carillo et al., *SIAM J. Applied Math.*, 48:1073 (1988).

Preferred methods to determine identity and/or similarity are designed to give the largest match between the sequences tested. Methods to determine identity and similarity are described in publicly available computer programs. Preferred computer program methods to determine identity and similarity between two sequences include, but are not limited to, the GCG program package, including GAP (Devereux et al., *Nucl. Acid. Res.*, 12:387 (1984); Genetics Computer Group, University of Wisconsin, Madison, WI), BLASTP, BLASTN, and FASTA (Altschul et al., *J. Mol. Biol.*, 215:403-410 (1990)). The BLASTX

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program is publicly available from the National Center for Biotechnology Information (NCBI) and other sources (*BLAST Manual*, Altschul et al. NCB/NLM/NIH Bethesda, MD 20894; Altschul et al., *supra*). The well known Smith Waterman  
5 algorithm may also be used to determine identity.

Certain alignment schemes for aligning two amino acid sequences may result in the matching of only a short region of the two sequences, and this small aligned region may have very high sequence identity even though there is no significant  
10 relationship between the two full length sequences. Accordingly, in a preferred embodiment, the selected alignment method (GAP program) will result in an alignment that spans at least 50 contiguous amino acids of the target polypeptide.

For example, using the computer algorithm GAP (Genetics  
15 Computer Group, University of Wisconsin, Madison, WI), two polypeptides for which the percent sequence identity is to be determined are aligned for optimal matching of their respective amino acids (the "matched span", as determined by the algorithm). A gap opening penalty (which is calculated as  
20 3X the average diagonal; the "average diagonal" is the average of the diagonal of the comparison matrix being used; the "diagonal" is the score or number assigned to each perfect amino acid match by the particular comparison matrix) and a gap extension penalty (which is usually 1/10 times the gap  
25 opening penalty), as well as a comparison matrix such as PAM 250 or BLOSUM 62 are used in conjunction with the algorithm. A standard comparison matrix (see Dayhoff et al., Atlas of Protein Sequence and Structure, vol. 5, supp.3 (1978) for the PAM 250 comparison matrix; Henikoff et al., Proc.  
30 Natl. Acad. Sci USA, 89:10915-10919 (1992) for the BLOSUM 62 comparison matrix) is also used by the algorithm.

Preferred parameters for a polypeptide sequence comparison include the following:

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Algorithm: Needleman et al., *J. Mol. Biol.*, 48:443-453 (1970);

Comparison matrix: BLOSUM 62 from Henikoff et al., *Proc. Natl. Acad. Sci. USA*, 89:10915-10919 (1992);

Gap Penalty: 12

Gap Length Penalty: 4

Threshold of Similarity: 0

10       The GAP program is useful with the above parameters. The aforementioned parameters are the default parameters for polypeptide comparisons (along with no penalty for end gaps) using the GAP algorithm.

15       Preferred parameters for nucleic acid molecule sequence comparisons include the following:

Algorithm: Needleman et al., *J. Mol Biol.*, 48:443-453 (1970);

Comparison matrix: matches = +10, mismatch = 0

Gap Penalty: 50

20       Gap Length Penalty: 3

The GAP program is also useful with the above parameters. The aforementioned parameters are the default parameters for nucleic acid molecule comparisons.

25       Other exemplary algorithms, gap opening penalties, gap extension penalties, comparison matrices, thresholds of similarity, etc. may be used,, including those set forth in the Program Manual, Wisconsin Package, Version 9, September, 1997. The particular choices to be made will be apparent to  
30 those of skill in the art and will depend on the specific comparison to be made, such as DNA to DNA, protein to protein, protein to DNA; and additionally, whether the comparison is

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between given pairs of sequences (in which case GAP or BestFit are generally preferred) or between one sequence and a large database of sequences (in which case FASTA or BLASTA are preferred).

5

### Synthesis

It will be appreciated by those skilled in the art the nucleic acid and polypeptide molecules described herein may be produced by recombinant and other means.

10

### Nucleic Acid Molecules

The nucleic acid molecules encode a polypeptide comprising the amino acid sequence of an IL-17 receptor like polypeptide can readily be obtained in a variety of ways including, without limitation, chemical synthesis, cDNA or genomic library screening, expression library screening and/or PCR amplification of cDNA.

Recombinant DNA methods used herein are generally those set forth in Sambrook et al., *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY (1989), and/or Ausubel et al., eds., *Current Protocols in Molecular Biology*, Green Publishers Inc. and Wiley and Sons, NY (1994). The present invention provides for nucleic acid molecules as described herein and methods for obtaining the molecules.

A gene or cDNA encoding a IL-17 receptor like polypeptide or fragment thereof may be obtained by hybridization screening of a genomic or cDNA library, or by PCR amplification. Where a gene encoding the amino acid sequence of an IL-17 receptor like polypeptide has been identified from one species, all or a portion of that gene may be used as a probe to identify orthologs or related genes from the same species. The probes or primers may be used to screen cDNA libraries from various

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tissue sources believed to express the IL-17 receptor like polypeptide. In addition, part or all of a nucleic acid molecule having the sequence as set forth in SEQ ID NO: 1 may be used to screen a genomic library to identify and isolate a gene encoding the amino acid sequence of an IL-17 receptor like polypeptide. Typically, conditions of moderate or high stringency will be employed for screening to minimize the number of false positives obtained from the screen.

Nucleic acid molecules encoding the amino acid sequence of IL-17 receptor like polypeptides may also be identified by expression cloning which employs the detection of positive clones based upon a property of the expressed protein. Typically, nucleic acid libraries are screened by the binding of an antibody or other binding partner (e.g., receptor or ligand) to cloned proteins which are expressed and displayed on a host cell surface. The antibody or binding partner is modified with a detectable label to identify those cells expressing the desired clone.

Recombinant expression techniques conducted in accordance with the descriptions set forth below may be followed to produce these polynucleotides and to express the encoded polypeptides. For example, by inserting a nucleic acid sequence which encodes the amino acid sequence of an IL-17 receptor like polypeptide into an appropriate vector, one skilled in the art can readily produce large quantities of the desired nucleotide sequence. The sequences can then be used to generate detection probes or amplification primers. Alternatively, a polynucleotide encoding the amino acid sequence of an IL-17 receptor like polypeptide can be inserted into an expression vector. By introducing the expression vector into an appropriate host, the encoded IL-17 receptor like polypeptide may be produced in large amounts.

Another method for obtaining a suitable nucleic acid sequence is the polymerase chain reaction (PCR). In this

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method, cDNA is prepared from poly(A)+RNA or total RNA using the enzyme reverse transcriptase. Two primers, typically complementary to two separate regions of cDNA (oligonucleotides) encoding the amino acid sequence of an IL-17 receptor like polypeptide, are then added to the cDNA along with a polymerase such as Tag polymerase, and the polymerase amplifies the cDNA region between the two primers.

Another means of preparing a nucleic acid molecule encoding the amino acid sequence of an IL-17 receptor like polypeptide, including a fragment or variant, is chemical synthesis using methods well known to the skilled artisan such as those described by Engels et al., *Angew. Chem. Intl. Ed.*, 28:716-734 (1989). These methods include, inter alia, the phosphotriester, phosphoramidite, and H-phosphonate methods for nucleic acid synthesis. A preferred method for such chemical synthesis is polymer-supported synthesis using standard phosphoramidite chemistry. Typically, the DNA encoding the amino acid sequence of an IL-17 receptor like polypeptide will be several hundred nucleotides in length. Nucleic acids larger than about 100 nucleotides can be synthesized as several fragments using these methods. The fragments can then be ligated together to form the full length nucleotide sequence of an IL-17 receptor like polypeptide. Usually, the DNA fragment encoding the amino terminus of the polypeptide will have an ATG, which encodes a methionine residue. This methionine may or may not be present on the mature form of the IL-17 receptor like polypeptide, depending on whether the polypeptide produced in the host cell is designed to be secreted from that cell. Other methods known to the skilled artisan may be used as well.

In some cases, it may be desirable to prepare nucleic acid molecules encoding IL-17 receptor like polypeptide variants. Nucleic acid molecules encoding variants may be produced using site directed mutagenesis, PCR amplification, or other appropriate methods, where the primer(s) have the desired point

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mutations (see Sambrook et al., supra, and Ausubel et al., supra, for descriptions of mutagenesis techniques). Chemical synthesis using methods described by Engels et al., supra, may also be used to prepare such variants. Other methods known to the skilled artisan may be used as well.

In certain embodiments, nucleic acid variants contain codons which have been altered for the optimal expression of an IL-17 receptor like polypeptide in a given host cell. Particular codon alterations will depend upon the IL-17 receptor like polypeptide(s) and host cell(s) selected for expression. Such "codon optimization" can be carried out by a variety of methods, for example, by selecting codons which are preferred for use in highly expressed genes in a given host cell. Computer algorithms which incorporate codon frequency tables such as "Ecohigh.cod" for codon preference of highly expressed bacterial genes may be used and are provided by the University of Wisconsin Package Version 9.0, Genetics Computer Group, Madison, WI. Other useful codon frequency tables include "Celegans\_high.cod", "Celegans\_low.cod", "Drosophila\_high.cod", "Human\_high.cod", "Maize\_high.cod", and "Yeast\_high.cod".

In other embodiments, nucleic acid molecules encode IL-17 receptor like variants with conservative amino acid substitutions as described herein, IL-17 receptor like variants comprising an addition and/or a deletion of one or more N-linked or O-linked glycosylation sites, IL-17 receptor like variants having deletions and/or substitutions of one or more cysteine residues, or IL-17 receptor like polypeptide fragments as described herein. In addition, nucleic acid molecules may encode any combination of IL-17 receptor like variants, fragments, and fusion polypeptides described herein.

### Vectors and Host Cells

A nucleic acid molecule encoding the amino acid sequence of an IL-17 receptor like polypeptide may be inserted into an



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appropriate expression vector using standard ligation techniques. The vector is typically selected to be functional in the particular host cell employed (i.e., the vector is compatible with the host cell machinery such that amplification of the gene and/or expression of the gene can occur). A nucleic acid molecule encoding the amino acid sequence of an IL-17 receptor like polypeptide may be amplified/expressed in prokaryotic, yeast, insect (baculovirus systems), and/or eukaryotic host cells. Selection of the host cell will depend in part on whether an IL-17 receptor like polypeptide is to be post-translationally modified (e.g., glycosylated and/or phosphorylated). If so, yeast, insect, or mammalian host cells are preferable. For a review of expression vectors, see *Meth. Enz.*, v.185, D.V. Goeddel, ed. Academic Press Inc., San Diego, CA (1990).

Typically, expression vectors used in any of the host cells will contain sequences for plasmid maintenance and for cloning and expression of exogenous nucleotide sequences. Such sequences, collectively referred to as "flanking sequences" in certain embodiments will typically include one or more of the following nucleotide sequences: a promoter, one or more enhancer sequences, an origin of replication, a transcriptional termination sequence, a complete intron sequence containing a donor and acceptor splice site, a sequence encoding a leader sequence for polypeptide secretion, a ribosome binding site, a polyadenylation sequence, a polylinker region for inserting the nucleic acid encoding the polypeptide to be expressed, and a selectable marker element. Each of these sequences is discussed below.

Optionally, the vector may contain a "tag"-encoding sequence, i.e., an oligonucleotide molecule located at the 5' or 3' end of the IL-17 receptor like polypeptide coding sequence; the oligonucleotide sequence encodes polyHis (such as hexaHis), or other "tag" such as FLAG, HA (hemagglutinin Influenza virus) or myc for which commercially available

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antibodies exist. This tag is typically fused to the polypeptide upon expression of the polypeptide, and can serve as a means for affinity purification of the IL-17 receptor like polypeptide from the host cell. Affinity purification can be accomplished, for example, by column chromatography using antibodies against the tag as an affinity matrix. Optionally, the tag can subsequently be removed from the purified IL-17 receptor like polypeptide by various means such as using certain peptidases for cleavage.

10 Flanking sequences may be homologous (i.e., from the same species and/or strain as the host cell), heterologous (i.e., from a species other than the host cell species or strain), hybrid (i.e., a combination of flanking sequences from more than one source) or synthetic, or the flanking sequences may  
15 be native sequences which normally function to regulate IL-17 receptor like polypeptide expression. As such, the source of a flanking sequence may be any prokaryotic or eukaryotic organism, any vertebrate or invertebrate organism, or any plant, provided that the flanking sequence is functional in,  
20 and can be activated by, the host cell machinery.

The flanking sequences useful in the vectors of this invention may be obtained by any of several methods well known in the art. Typically, flanking sequences useful herein other than the IL-17 receptor like gene flanking sequences will have  
25 been previously identified by mapping and/or by restriction endonuclease digestion and can thus be isolated from the proper tissue source using the appropriate restriction endonucleases. In some cases, the full nucleotide sequence of a flanking sequence may be known. Here, the flanking sequence  
30 may be synthesized using the methods described herein for nucleic acid synthesis or cloning.

Where all or only a portion of the flanking sequence is known, it may be obtained using PCR and/or by screening a genomic library with suitable oligonucleotide and/or flanking

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sequence fragments from the same or another species. Where the flanking sequence is not known, a fragment of DNA containing a flanking sequence may be isolated from a larger piece of DNA that may contain, for example, a coding sequence or even another gene or genes. Isolation may be accomplished by restriction endonuclease digestion to produce the proper DNA fragment followed by isolation using agarose gel purification, Qiagen® column chromatography (Chatsworth, CA), or other methods known to the skilled artisan. The selection of suitable enzymes to accomplish this purpose will be readily apparent to one of ordinary skill in the art.

An origin of replication is typically a part of those prokaryotic expression vectors purchased commercially, and the origin aids in the amplification of the vector in a host cell. Amplification of the vector to a certain copy number can, in some cases, be important for the optimal expression of an IL-17 receptor like polypeptide. If the vector of choice does not contain an origin of replication site, one may be chemically synthesized based on a known sequence, and ligated into the vector. For example, the origin of replication from the plasmid pBR322 (Product No. 303-3s, New England Biolabs, Beverly, MA) is suitable for most Gram-negative bacteria and various origins (e.g., SV40, polyoma, adenovirus, vesicular stomatitis virus (VSV) or papillomaviruses such as HPV or BPV) are useful for cloning vectors in mammalian cells. Generally, the origin of replication component is not needed for mammalian expression vectors (for example, the SV40 origin is often used only because it contains the early promoter).

A transcription termination sequence is typically located 3' of the end of a polypeptide coding region and serves to terminate transcription. Usually, a transcription termination sequence in prokaryotic cells is a G-C rich fragment followed by a poly T sequence. While the sequence is easily cloned from a library or even purchased commercially as part of a

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vector, it can also be readily synthesized using methods for nucleic acid synthesis such as those described herein.

A selectable marker gene element encodes a protein necessary for the survival and growth of a host cell grown in a selective culture medium. Typical selection marker genes encode proteins that (a) confer resistance to antibiotics or other toxins, e.g., ampicillin, tetracycline, or kanamycin for prokaryotic host cells, (b) complement auxotrophic deficiencies of the cell; or (c) supply critical nutrients not available from complex media. Preferred selectable markers are the kanamycin resistance gene, the ampicillin resistance gene, and the tetracycline resistance gene. A neomycin resistance gene may also be used for selection in prokaryotic and eukaryotic host cells.

Other selection genes may be used to amplify the gene which will be expressed. Amplification is the process wherein genes which are in greater demand for the production of a protein critical for growth are reiterated in tandem within the chromosomes of successive generations of recombinant cells. Examples of suitable selectable markers for mammalian cells include dihydrofolate reductase (DHFR) and thymidine kinase. The mammalian cell transformants are placed under selection pressure which only the transformants are uniquely adapted to survive by virtue of the selection gene present in the vector. Selection pressure is imposed by culturing the transformed cells under conditions in which the concentration of selection agent in the medium is successively changed, thereby leading to the amplification of both the selection gene and the DNA that encodes an IL-17 receptor like polypeptide. As a result, increased quantities of IL-17 receptor like polypeptide are synthesized from the amplified DNA.

A ribosome binding site is usually necessary for translation initiation of mRNA and is characterized by a

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Shine-Dalgarno sequence (prokaryotes) or a Kozak sequence (eukaryotes). The element is typically located 3' to the promoter and 5' to the coding sequence of an IL-17 receptor like polypeptide to be expressed. The Shine-Dalgarno sequence is varied but is typically a polypurine (i.e., having a high A-G content). Many Shine-Dalgarno sequences have been identified, each of which can be readily synthesized using methods set forth herein and used in a prokaryotic vector.

A leader, or signal, sequence may be used to direct an IL-17 receptor like polypeptide out of the host cell. Typically, a nucleotide sequence encoding the signal sequence is positioned in the coding region of an IL-17 receptor like nucleic acid molecule, or directly at the 5' end of an IL-17 receptor like polypeptide coding region. Many signal sequences have been identified, and any of those that are functional in the selected host cell may be used in conjunction with an IL-17 receptor like nucleic acid molecule. Therefore, a signal sequence may be homologous (naturally occurring) or heterologous to an IL-17 receptor like gene or cDNA. Additionally, a signal sequence may be chemically synthesized using methods described herein. In most cases, the secretion of an IL-17 receptor like polypeptide from the host cell via the presence of a signal peptide will result in the removal of the signal peptide from the secreted IL-17 receptor like polypeptide. The signal sequence may be a component of the vector, or it may be a part of an IL-17 receptor like nucleic acid molecule that is inserted into the vector.

Included within the scope of this invention is the use of either a nucleotide sequence encoding a native IL-17 receptor like polypeptide signal sequence joined to an IL-17 receptor like polypeptide coding region or a nucleotide sequence encoding a heterologous signal sequence joined to an IL-17 receptor like polypeptide coding region. The heterologous signal sequence selected should be one that is recognized and

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processed, i.e., cleaved by a signal peptidase, by the host cell. For prokaryotic host cells that do not recognize and process the native IL-17 receptor like polypeptide signal sequence, the signal sequence is substituted by a prokaryotic signal sequence selected, for example, from the group of the alkaline phosphatase, penicillinase, or heat-stable enterotoxin II leaders. For yeast secretion, the native IL-17 receptor like polypeptide signal sequence may be substituted by the yeast invertase, alpha factor, or acid phosphatase leaders. In mammalian cell expression the native signal sequence is satisfactory, although other mammalian signal sequences may be suitable.

In some cases, such as where glycosylation is desired in a eukaryotic host cell expression system, one may manipulate the various presequences to improve glycosylation or yield. For example, one may alter the peptidase cleavage site of a particular signal peptide, or add presequences, which also may affect glycosylation. The final protein product may have, in the -1 position (relative to the first amino acid of the mature protein) one or more additional amino acids incident to expression, which may not have been totally removed. For example, the final protein product may have one or two amino acid residues found in the peptidase cleavage site, attached to the N-terminus. Alternatively, use of some enzyme cleavage sites may result in a slightly truncated form of the desired IL-17 receptor like polypeptide, if the enzyme cuts at such area within the mature polypeptide.

In many cases, transcription of a nucleic acid molecule is increased by the presence of one or more introns in the vector; this is particularly true where a polypeptide is produced in eukaryotic host cells, especially mammalian host cells. The introns used may be naturally occurring within the IL-17 receptor like gene, especially where the gene used is a full length genomic sequence or a fragment thereof. Where the intron is not naturally occurring within the gene (as for most

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cDNAs), the intron(s) may be obtained from another source. The position of the intron with respect to flanking sequences and the IL-17 receptor like gene is generally important, as the intron must be transcribed to be effective. Thus, when an IL-17 receptor like cDNA molecule is being transcribed, the preferred position for the intron is 3' to the transcription start site, and 5' to the polyA transcription termination sequence. Preferably, the intron or introns will be located on one side or the other (i.e., 5' or 3') of the cDNA such that it does not interrupt the coding sequence. Any intron from any source, including any viral, prokaryotic and eukaryotic (plant or animal) organisms, may be used to practice this invention, provided that it is compatible with the host cell(s) into which it is inserted. Also included herein are synthetic introns. Optionally, more than one intron may be used in the vector.

The expression and cloning vectors of the present invention will each typically contain a promoter that is recognized by the host organism and operably linked to the molecule encoding a IL-17 receptor like polypeptide. Promoters are untranscribed sequences located upstream (5') to the start codon of a structural gene (generally within about 100 to 1000 bp) that control the transcription of the structural gene. Promoters are conventionally grouped into one of two classes, inducible promoters and constitutive promoters. Inducible promoters initiate increased levels of transcription from DNA under their control in response to some change in culture conditions, such as the presence or absence of a nutrient or a change in temperature. Constitutive promoters, on the other hand, initiate continual gene product production; that is, there is little or no control over gene expression. A large number of promoters, recognized by a variety of potential host cells, are well known. A suitable promoter is operably linked to the DNA encoding an IL-17 receptor like polypeptide by removing the promoter from the

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source DNA by restriction enzyme digestion and inserting the desired promoter sequence into the vector. The native IL-17 receptor like gene promoter sequence may be used to direct amplification and/or expression of an IL-17 receptor like nucleic acid molecule. A heterologous promoter is preferred, however, if it permits greater transcription and higher yields of the expressed protein as compared to the native promoter, and if it is compatible with the host cell system that has been selected for use.

10 Promoters suitable for use with prokaryotic hosts include the beta-lactamase and lactose promoter systems; alkaline phosphatase, a tryptophan (trp) promoter system; and hybrid promoters such as the tac promoter. Other known bacterial promoters are also suitable. Their sequences have been  
15 published, thereby enabling one skilled in the art to ligate them to the desired DNA sequence(s), using linkers or adapters as needed to supply any useful restriction sites.

Suitable promoters for use with yeast hosts are also well known in the art. Yeast enhancers are advantageously used  
20 with yeast promoters. Suitable promoters for use with mammalian host cells are well known and include, but are not limited to, those obtained from the genomes of viruses such as polyoma virus, fowlpox virus, adenovirus (such as Adenovirus 2), bovine papilloma virus, avian sarcoma virus,  
25 cytomegalovirus (CMV), a retrovirus, hepatitis-B virus and most preferably Simian Virus 40 (SV40). Other suitable mammalian promoters include heterologous mammalian promoters, e.g., heat-shock promoters and the actin promoter.

Additional promoters which may be of interest in  
30 controlling IL-17 receptor like gene transcription include, but are not limited to: the SV40 early promoter region (Bernoist and Chambon, *Nature*, 290:304-310, 1981); the CMV promoter; the promoter contained in the 3' long terminal repeat of Rous sarcoma virus (Yamamoto et al., *Cell*, 22:787-



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797, 1980); the herpes thymidine kinase promoter (Wagner et al., *Proc. Natl. Acad. Sci. USA*, 78:144-1445, 1981); the regulatory sequences of the metallothionine gene (Brinster et al., *Nature*, 296:39-42, 1982); prokaryotic expression vectors  
5 such as the beta-lactamase promoter (Villa-Kamaroff, et al., *Proc. Natl. Acad. Sci. USA*, 75:3727-3731, 1978); or the tac promoter (DeBoer, et al., *Proc. Natl. Acad. Sci. USA*, 80:21-25, 1983). Also of interest are the following animal transcriptional control regions, which exhibit tissue  
10 specificity and have been utilized in transgenic animals: the elastase I gene control region which is active in pancreatic acinar cells (Swift et al., *Cell*, 38:639-646, 1984; Ornitz et al., *Cold Spring Harbor Symp. Quant. Biol.*, 50:399-409 (1986); MacDonald, *Hepatology*, 7:425-515, 1987); the insulin gene  
15 control region which is active in pancreatic beta cells (Hanahan, *Nature*, 315:115-122, 1985); the immunoglobulin gene control region which is active in lymphoid cells (Grosschedl et al., *Cell*, 38:647-658 (1984); Adames et al., *Nature*, 318:533-538 (1985); Alexander et al., *Mol. Cell. Biol.*, 7:1436-1444, 1987); the mouse mammary tumor virus control  
20 region which is active in testicular, breast, lymphoid and mast cells (Leder et al., *Cell*, 45:485-495, 1986); the albumin gene control region which is active in liver (Pinkert et al., *Genes and Devel.*, 1:268-276, 1987); the alphafetoprotein gene  
25 control region which is active in liver (Krumlauf et al., *Mol. Cell. Biol.*, 5:1639-1648, 1985; Hammer et al., *Science*, 235:53-58, 1987); the alpha 1-antitrypsin gene control region which is active in the liver (Kelsey et al., *Genes and Devel.*, 1:161-171, 1987); the beta-globin gene control region which is  
30 active in myeloid cells (Mogram et al., *Nature*, 315:338-340, 1985; Kollias et al., *Cell*, 46:89-94, 1986); the myelin basic protein gene control region which is active in oligodendrocyte cells in the brain (Readhead et al., *Cell*, 48:703-712, 1987); the myosin light chain-2 gene control region which is active  
35 in skeletal muscle (Sani, *Nature*, 314:283-286, 1985); and the

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gonadotropic releasing hormone gene control region which is active in the hypothalamus (Mason *et al.*, *Science*, 234:1372-1378, 1986).

An enhancer sequence may be inserted into the vector to  
5 increase the transcription of a DNA encoding an IL-17 receptor  
like polypeptide of the present invention by higher  
eukaryotes. Enhancers are cis-acting elements of DNA, usually  
about 10-300 bp in length, that act on the promoter to  
increase transcription. Enhancers are relatively orientation  
10 and position independent. They have been found 5' and 3' to  
the transcription unit. Several enhancer sequences available  
from mammalian genes are known (e.g., globin, elastase,  
albumin, alpha-feto-protein and insulin). Typically, however,  
an enhancer from a virus will be used. The SV40 enhancer, the  
15 cytomegalovirus early promoter enhancer, the polyoma enhancer,  
and adenovirus enhancers are exemplary enhancing elements for  
the activation of eukaryotic promoters. While an enhancer may  
be spliced into the vector at a position 5' or 3' to an IL-17  
RECEPTOR LIKE nucleic acid molecule, it is typically located  
20 at a site 5' from the promoter.

Expression vectors of the invention may be constructed  
from a starting vector such as a commercially available  
vector. Such vectors may or may not contain all of the  
desired flanking sequences. Where one or more of the desired  
25 flanking sequences are not already present in the vector, they  
may be individually obtained and ligated into the vector.  
Methods used for obtaining each of the flanking sequences are  
well known to one skilled in the art.

Preferred vectors for practicing this invention are those  
30 which are compatible with bacterial, insect, and mammalian  
host cells. Such vectors include, *inter alia*, pCRII, pCR3,  
and pCDNA3.1 (Invitrogen Company, Carlsbad, CA), pBSII  
(Stratagene Company, La Jolla, CA), pET15 (Novagen, Madison,  
WI), pGEX (Pharmacia Biotech, Piscataway, NJ), pEGFP-N2

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(Clontech, Palo Alto, CA), pETL (BlueBacII; Invitrogen), pDSR-alpha (PCT Publication No. WO90/14363) and pFastBacDual (Gibco/BRL, Grand Island, NY).

Additional suitable vectors include, but are not limited to, cosmids, plasmids or modified viruses, but it will be appreciated that the vector system must be compatible with the selected host cell. Such vectors include, but are not limited to plasmids such as Bluescript<sup>®</sup> plasmid derivatives (a high copy number ColE1-based phagemid, Stratagene Cloning Systems Inc., La Jolla CA), PCR cloning plasmids designed for cloning Taq-amplified PCR products (e.g., TOPO<sup>™</sup> TA Cloning<sup>®</sup> Kit, PCR2.1<sup>®</sup> plasmid derivatives, Invitrogen, Carlsbad, CA), and mammalian, yeast, or virus vectors such as a baculovirus expression system (pBacPAK plasmid derivatives, Clontech, Palo Alto, CA). The recombinant molecules can be introduced into host cells via transformation, transfection, infection, electroporation or other known techniques.

After the vector has been constructed and a nucleic acid molecule encoding an IL-17 receptor like polypeptide has been inserted into the proper site of the vector, the completed vector may be inserted into a suitable host cell for amplification and/or polypeptide expression. The transformation of an expression vector for an IL-17 receptor like polypeptide into a selected host cell may be accomplished by well known methods including methods such as transfection, infection, calcium chloride, electroporation, microinjection, lipofection or the DEAE-dextran method or other known techniques. The method selected will in part be a function of the type of host cell to be used. These methods and other suitable methods are well known to the skilled artisan, and are set forth, for example, in Sambrook et al., *supra*.

Host cells may be prokaryotic host cells (such as *E. coli*) or eukaryotic host cells (such as a yeast cell, an insect cell or a vertebrate cell). The host cell, when

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cultured under appropriate conditions, synthesizes an IL-17 receptor like polypeptide which can subsequently be collected from the culture medium (if the host cell secretes it into the medium) or directly from the host cell producing it (if it is not secreted). The selection of an appropriate host cell will depend upon various factors, such as desired expression levels, polypeptide modifications that are desirable or necessary for activity, such as glycosylation or phosphorylation, and ease of folding into a biologically active molecule.

A number of suitable host cells are known in the art and many are available from the American Type Culture Collection (ATCC), 10801 University Boulevard, Manassas, VA 20110-2209. Examples include, but are not limited to, mammalian cells, such as Chinese hamster ovary cells (CHO) (ATCC No. CCL61) CHO DHFR-cells (Urlaub et al., *Proc. Natl. Acad. Sci. USA*, 97:4216-4220 (1980)), human embryonic kidney (HEK) 293 or 293T cells (ATCC No. CRL1573), or 3T3 cells (ATCC No. CCL92). The selection of suitable mammalian host cells and methods for transformation, culture, amplification, screening and product production and purification are known in the art. Other suitable mammalian cell lines, are the monkey COS-1 (ATCC No. CRL1650) and COS-7 cell lines (ATCC No. CRL1651), and the CV-1 cell line (ATCC No. CCL70). Further exemplary mammalian host cells include primate cell lines and rodent cell lines, including transformed cell lines. Normal diploid cells, cell strains derived from *in vitro* culture of primary tissue, as well as primary explants, are also suitable. Candidate cells may be genotypically deficient in the selection gene, or may contain a dominantly acting selection gene. Other suitable mammalian cell lines include but are not limited to, mouse neuroblastoma N2A cells, HeLa, mouse L-929 cells, 3T3 lines derived from Swiss, Balb-c or NIH mice, BHK or HaK hamster cell lines, which are available from the ATCC. Each of these cell lines is known by and available to those skilled in the

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art of protein expression.

Similarly useful as host cells suitable for the present invention are bacterial cells. For example, the various strains of *E. coli* (e.g., HB101, (ATCC No. 33694) DH5 $\alpha$ , DH10, and MC1061 (ATCC No. 53338)) are well-known as host cells in the field of biotechnology. Various strains of *B. subtilis*, *Pseudomonas* spp., other *Bacillus* spp., *Streptomyces* spp., and the like may also be employed in this method.

Many strains of yeast cells known to those skilled in the art are also available as host cells for the expression of the polypeptides of the present invention. Preferred yeast cells include, for example, *Saccharomyces cerevisiae* and *Pichia pastoris*.

Additionally, where desired, insect cell systems may be utilized in the methods of the present invention. Such systems are described for example in Kitts et al., *Biotechniques*, 14:810-817 (1993); Lucklow, *Curr. Opin. Biotechnol.*, 4:564-572 (1993); and Lucklow et al. (*J. Virol.*, 67:4566-4579 (1993)). Preferred insect cells are Sf-9 and Hi5 (Invitrogen, Carlsbad, CA).

One may also use transgenic animals to express glycosylated IL-17 receptor like polypeptides. For example, one may use a transgenic milk-producing animal (a cow or goat, for example) and obtain the present glycosylated polypeptide in the animal milk. One may also use plants to produce IL-17 receptor like polypeptides, however, in general, the glycosylation occurring in plants is different from that produced in mammalian cells, and may result in a glycosylated product which is not suitable for human therapeutic use.

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Polypeptide Production

Host cells comprising an IL-17 receptor like polypeptide expression vector may be cultured using standard media well known to the skilled artisan. The media will usually contain all nutrients necessary for the growth and survival of the cells. Suitable media for culturing *E. coli* cells include, for example, Luria Broth (LB) and/or Terrific Broth (TB). Suitable media for culturing eukaryotic cells include Roswell Park Memorial Institute medium 1640 (RPMI 1640), Minimal Essential Medium (MEM) and/or Dulbecco's Modified Eagle Medium (DMEM), all of which may be supplemented with serum and/or growth factors as indicated by the particular cell line being cultured. A suitable medium for insect cultures is Grace's medium supplemented with yeastolate, lactalbumin hydrolysate and/or fetal calf serum, as necessary.

Typically, an antibiotic or other compound useful for selective growth of transformed cells is added as a supplement to the media. The compound to be used will be dictated by the selectable marker element present on the plasmid with which the host cell was transformed. For example, where the selectable marker element is kanamycin resistance, the compound added to the culture medium will be kanamycin. Other compounds for selective growth include ampicillin, tetracycline, and neomycin.

The amount of an IL-17 receptor like polypeptide produced by a host cell can be evaluated using standard methods known in the art. Such methods include, without limitation, Western blot analysis, SDS-polyacrylamide gel electrophoresis, non-denaturing gel electrophoresis, high performance liquid chromatography (HPLC) separation, immunoprecipitation, and/or activity assays such as DNA binding gel shift assays.

If an IL-17 receptor like polypeptide has been designed to be secreted from the host cells, the majority of polypeptide may be found in the cell culture medium. If

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however, the IL-17 receptor like polypeptide is not secreted from the host cells, it will be present in the cytoplasm and/or the nucleus (for eukaryotic host cells) or in the cytosol (for bacterial host cells).

5 For an IL-17 receptor like polypeptide situated in the host cell cytoplasm and/or the nucleus (for eukaryotic host cells) or in the cytosol (for bacterial host cells), intracellular material (including inclusion bodies for gram-negative bacteria) can be extracted from the host cell using  
10 any standard technique known to the skilled artisan. For example, the host cells can be lysed to release the contents of the periplasm/cytoplasm by French press, homogenization, and/or sonication followed by centrifugation.

If an IL-17 receptor like polypeptide has formed  
15 inclusion bodies in the cytosol, the inclusion bodies can often bind to the inner and/or outer cellular membranes and thus will be found primarily in the pellet material after centrifugation. The pellet material can then be treated at pH extremes or with a chaotropic agent such as a detergent,  
20 guanidine, guanidine derivatives, urea, or urea derivatives in the presence of a reducing agent such as dithiothreitol at alkaline pH or tris carboxyethyl phosphine at acid pH to release, break apart, and solubilize the inclusion bodies. The IL-17 receptor like polypeptide in its now soluble form  
25 can then be analyzed using gel electrophoresis, immunoprecipitation or the like. If it is desired to isolate the IL-17 receptor like polypeptide, isolation may be accomplished using standard methods such as those described herein and in Marston et al., *Meth. Enz.*, 182:264-275 (1990).

30 In some cases, an IL-17 receptor like polypeptide may not be biologically active upon isolation. Various methods for "refolding" or converting the polypeptide to its tertiary structure and generating disulfide linkages can be used to restore biological activity. Such methods include exposing

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the solubilized polypeptide to a pH usually above 7 and in the presence of a particular concentration of a chaotrope. The selection of chaotrope is very similar to the choices used for inclusion body solubilization, but usually the chaotrope is used at a lower concentration and is not necessarily the same as chaotropes used for the solubilization. In most cases the refolding/oxidation solution will also contain a reducing agent or the reducing agent plus its oxidized form in a specific ratio to generate a particular redox potential allowing for disulfide shuffling to occur in the formation of the protein's cysteine bridge(s). Some of the commonly used redox couples include cysteine/cystamine, glutathione (GSH)/dithiobis GSH, cupric chloride, dithiothreitol (DTT)/dithiane DTT, and 2-mercaptoethanol (BME)/dithio-B (ME). A cosolvent may be used to increase the efficiency of the refolding, and the more common reagents used for this purpose include glycerol, polyethylene glycol of various molecular weights, arginine and the like.

If inclusion bodies are not formed to a significant degree upon expression of an IL-17 receptor like polypeptide, then the polypeptide will be found primarily in the supernatant after centrifugation of the cell homogenate. The polypeptide may be further isolated from the supernatant using methods such as those described herein.

The purification of an IL-17 receptor like polypeptide from solution can be accomplished using a variety of techniques. If the polypeptide has been synthesized such that it contains a tag such as Hexahistidine (IL-17 receptor like polypeptide/hexaHis) or other small peptide such as FLAG (Eastman Kodak Co., New Haven, CT) or myc (Invitrogen, Carlsbad, CA) at either its carboxyl or amino terminus, it may be purified in a one-step process by passing the solution through an affinity column where the column matrix has a high affinity for the tag.



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For example, polyhistidine binds with great affinity and specificity to nickel, thus an affinity column of nickel (such as the Qiagen® nickel columns) can be used for purification of IL-17 receptor like polypeptide/polyHis. See for example, 5 Ausubel et al., eds., *Current Protocols in Molecular Biology*, Section 10.11.8, John Wiley & Sons, New York (1993).

Additionally, the IL-17 receptor like polypeptide may be purified through the use of a monoclonal antibody which is capable of specifically recognizing and binding to the IL-17 10 receptor like polypeptide.

Suitable procedures for purification thus include, without limitation, affinity chromatography, immunoaffinity chromatography, ion exchange chromatography, molecular sieve chromatography, High Performance Liquid Chromatography (HPLC), 15 electrophoresis (including native gel electrophoresis) followed by gel elution, and preparative isoelectric focusing ("Isoprime" machine/technique, Hoefer Scientific, San Francisco, CA). In some cases, two or more purification techniques may be combined to achieve increased purity.

20 IL-17 receptor like polypeptides, including fragments, variants, and/or derivatives thereof may also be prepared by chemical synthesis methods (such as solid phase peptide synthesis) using techniques known in the art, such as those set forth by Merrifield et al., *J. Am. Chem. Soc.*, 85:2149 25 (1963), Houghten et al., *Proc Natl Acad. Sci. USA*, 82:5132 (1985), and Stewart and Young, *Solid Phase Peptide Synthesis*, Pierce Chemical Co., Rockford, IL (1984). Such polypeptides may be synthesized with or without a methionine on the amino terminus. Chemically synthesized IL-17 receptor like 30 polypeptides may be oxidized using methods set forth in these references to form disulfide bridges. Chemically synthesized IL-17 receptor like polypeptides are expected to have comparable biological activity to the corresponding IL-17

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receptor like polypeptides produced recombinantly or purified from natural sources, and thus may be used interchangeably with a recombinant or natural IL-17 receptor like polypeptide.

Another means of obtaining an IL-17 receptor like polypeptide is via purification from biological samples such as source tissues and/or fluids in which the IL-17 receptor like polypeptide is naturally found. Such purification can be conducted using methods for protein purification as described herein. The presence of the IL-17 receptor like polypeptide during purification may be monitored using, for example, an antibody prepared against recombinantly produced IL-17 receptor like polypeptide or peptide fragments thereof.

A number of additional methods for producing nucleic acids and polypeptides are known in the art, and can be used to produce polypeptides having specificity for IL-17 receptor like. See for example, Roberts et al., *Proc. Natl. Acad. Sci.*, 94:12297-12303 (1997), which describes the production of fusion proteins between an mRNA and its encoded peptide. See also Roberts, R., *Curr. Opin. Chem. Biol.*, 3:268-273 (1999). Additionally, U.S. patent No. 5,824,469 describes methods of obtaining oligonucleotides capable of carrying out a specific biological function. The procedure involves generating a heterogeneous pool of oligonucleotides, each having a 5' randomized sequence, a central preselected sequence, and a 3' randomized sequence. The resulting heterogeneous pool is introduced into a population of cells that do not exhibit the desired biological function. Subpopulations of the cells are then screened for those which exhibit a predetermined biological function. From that subpopulation, oligonucleotides capable of carrying out the desired biological function are isolated.

U.S. Patent Nos. 5,763,192, 5,814,476, 5,723,323, and 5,817,483 describe processes for producing peptides or polypeptides. This is done by producing stochastic genes or

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fragments thereof, and then introducing these genes into host cells which produce one or more proteins encoded by the stochastic genes. The host cells are then screened to identify those clones producing peptides or polypeptides  
5 having the desired activity.

Another method for producing peptides or polypeptides is described in PCT/US98/20094 (WO99/15650) filed by Athersys, Inc. Known as "Random Activation of Gene Expression for Gene Discovery" (RAGE-GD), the process involves  
10 the activation of endogenous gene expression or over-expression of a gene by in situ recombination methods. For example, expression of an endogenous gene is activated or increased by integrating a regulatory sequence into the target cell which is capable of activating expression of the gene by  
15 non-homologous or illegitimate recombination. The target DNA is first subjected to radiation, and a genetic promoter inserted. The promoter eventually locates a break at the front of a gene, initiating transcription of the gene. This results in expression of the desired peptide or polypeptide.

20 It will be appreciated that these methods can also be used to create comprehensive IL-17 RECEPTOR like protein expression libraries, which can subsequently be used for high throughput phenotypic screening in a variety of assays, such as biochemical assays, cellular assays, and whole organism  
25 assays (e.g., plant, mouse, etc.).

### Chemical Derivatives

Chemically modified derivatives of the IL-17 receptor like polypeptides may be prepared by one skilled in the art, given the disclosures set forth hereinbelow. IL-17 receptor  
30 like polypeptide derivatives are modified in a manner that is different, either in the type or location of the molecules naturally attached to the polypeptide. Derivatives may include molecules formed by the deletion of one or more naturally-attached chemical groups. The polypeptide

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comprising the amino acid sequence of SEQ ID NO: 2, or an IL-17 receptor like polypeptide variant may be modified by the covalent attachment of one or more polymers. For example, the polymer selected is typically water soluble so that the protein to which it is attached does not precipitate in an aqueous environment, such as a physiological environment. Included within the scope of suitable polymers is a mixture of polymers. Preferably, for therapeutic use of the end-product preparation, the polymer will be pharmaceutically acceptable.

The polymers each may be of any molecular weight and may be branched or unbranched. The polymers each typically have an average molecular weight of between about 2kDa to about 100kDa (the term "about" indicating that in preparations of a water soluble polymer, some molecules will weigh more, some less, than the stated molecular weight). The average molecular weight of each polymer preferably is between about 5kDa and about 50kDa, more preferably between about 12kDa and about 40kDa and most preferably between about 20kDa and about 35kDa.

Suitable water soluble polymers or mixtures thereof include, but are not limited to, N-linked or O-linked carbohydrates, sugars, phosphates, polyethylene glycol (PEG) (including the forms of PEG that have been used to derivatize proteins, including mono-(C<sub>1</sub>-C<sub>10</sub>) alkoxy- or aryloxy-polyethylene glycol), monomethoxy-polyethylene glycol, dextran (such as low molecular weight dextran, of, for example about 6 kD), cellulose, or other carbohydrate based polymers, poly-(N-vinyl pyrrolidone) polyethylene glycol, propylene glycol homopolymers, a polypropylene oxide/ethylene oxide co-polymer, polyoxyethylated polyols (e.g., glycerol) and polyvinyl alcohol. Also encompassed by the present invention are bifunctional crosslinking molecules which may be used to prepare covalently attached multimers of the polypeptide

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comprising the amino acid sequence of SEQ ID NO: 2 or an IL-17 receptor like polypeptide variant.

In general, chemical derivatization may be performed under any suitable condition used to react a protein with an activated polymer molecule. Methods for preparing chemical derivatives of polypeptides will generally comprise the steps of (a) reacting the polypeptide with the activated polymer molecule (such as a reactive ester or aldehyde derivative of the polymer molecule) under conditions whereby the polypeptide comprising the amino acid sequence of SEQ ID NO: 2, or an IL-17 receptor like polypeptide variant becomes attached to one or more polymer molecules, and (b) obtaining the reaction product(s). The optimal reaction conditions will be determined based on known parameters and the desired result. For example, the larger the ratio of polymer molecules:protein, the greater the percentage of attached polymer molecule. In one embodiment, the IL-17 receptor like polypeptide derivative may have a single polymer molecule moiety at the amino terminus. See, for example, U.S. Patent No. 5,234,784.

The pegylation of the polypeptide specifically may be carried out by any of the pegylation reactions known in the art, as described for example in the following references: Francis et al., *Focus on Growth Factors*, 3:4-10 (1992); EP 0154316; EP 0401384 and U.S. Patent No. 4,179,337. For example, pegylation may be carried out via an acylation reaction or an alkylation reaction with a reactive polyethylene glycol molecule (or an analogous reactive water-soluble polymer) as described herein. For the acylation reactions, the polymer(s) selected should have a single reactive ester group. For reductive alkylation, the polymer(s) selected should have a single reactive aldehyde group. A reactive aldehyde is, for example, polyethylene glycol propionaldehyde, which is water stable, or mono C<sub>1</sub>-C<sub>10</sub>

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alkoxy or aryloxy derivatives thereof (see U.S. Patent No. 5,252,714).

In another embodiment, IL-17 receptor like polypeptides may be chemically coupled to biotin, and the biotin/IL-17  
5 receptor like polypeptide molecules which are conjugated are then allowed to bind to avidin, resulting in tetravalent avidin/biotin/IL-17 receptor like polypeptide molecules. IL-17 receptor like polypeptides may also be covalently coupled to dinitrophenol (DNP) or trinitrophenol (TNP) and the  
10 resulting conjugates precipitated with anti-DNP or anti-TNP-IgM to form decameric conjugates with a valency of 10.

Generally, conditions which may be alleviated or modulated by the administration of the present IL-17 receptor like polypeptide derivatives include those described herein  
15 for IL-17 receptor like polypeptides. However, the IL-17 receptor like polypeptide derivatives disclosed herein may have additional activities, enhanced or reduced biological activity, or other characteristics, such as increased or decreased half-life, as compared to the non-derivatized  
20 molecules.

The present invention further includes non-human animals in which the promoter for one or more of the IL-17 receptor like polypeptides of the present invention is either activated or inactivated (e.g., by using homologous recombination  
25 methods) to alter the level of expression of one or more of the native IL-17 receptor like polypeptides.

These non-human animals may be used for drug candidate screening. In such screening, the impact of a drug candidate  
30 on the animal may be measured. For example, drug candidates may decrease or increase the expression of the IL-17 receptor like gene. In certain embodiments, the amount of IL-17 receptor like polypeptide, that is produced may be measured

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after the exposure of the animal to the drug candidate. Additionally, in certain embodiments, one may detect the actual impact of the drug candidate on the animal. For example, the overexpression of a particular gene may result in, or be associated with, a disease or pathological condition. In such cases, one may test a drug candidate's ability to decrease expression of the gene or its ability to prevent or inhibit a pathological condition. In other examples, the production of a particular metabolic product such as a fragment of a polypeptide, may result in, or be associated with, a disease or pathological condition. In such cases, one may test a drug candidate's ability to decrease the production of such a metabolic product or its ability to prevent or inhibit a pathological condition.

15

#### Microarray

It will be appreciated that DNA microarray technology can be utilized in accordance with the present invention. DNA microarrays are miniature, high density arrays of nucleic acids positioned on a solid support, such as glass. Each cell or element within the array has numerous copies of a single species of DNA which acts as a target for hybridization for its cognate mRNA. In expression profiling using DNA microarray technology, mRNA is first extracted from a cell or tissue sample and then converted enzymatically to fluorescently labeled cDNA. This material is hybridized to the microarray and unbound cDNA is removed by washing. The expression of discrete genes represented on the array is then visualized by quantitating the amount of labeled cDNA which is specifically bound to each target DNA. In this way, the expression of thousands of genes can be quantitated in a high throughput, parallel manner from a single sample of biological material.

This high throughput expression profiling has a broad range of applications with respect to the IL-17 receptor like

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molecules of the invention, including, but not limited to: the identification and validation of IL-17 receptor like disease-related genes as targets for therapeutics; molecular toxicology of IL-17 receptor like molecules and inhibitors thereof; stratification of populations and generation of surrogate markers for clinical trials; and enhancing IL-17 receptor like-related small molecule drug discovery by aiding in the identification of selective compounds in high throughput screens (HTS).

#### 10 Selective Binding Agents

As used herein, the term "selective binding agent" refers to a molecule which has specificity for one or more IL-17 receptor like polypeptides. Suitable selective binding agents include, but are not limited to, antibodies and derivatives thereof, polypeptides, and small molecules. Suitable selective binding agents may be prepared using methods known in the art. An exemplary IL-17 receptor like polypeptide selective binding agent of the present invention is capable of binding a certain portion of the IL-17 receptor like polypeptide thereby inhibiting the binding of the polypeptide to the IL-17 receptor like polypeptide receptor(s).

Selective binding agents such as antibodies and antibody fragments that bind IL-17 receptor like polypeptides are within the scope of the present invention. The antibodies may be polyclonal including monospecific polyclonal, monoclonal (MAbs), recombinant, chimeric, humanized such as CDR-grafted, human, single chain, and/or bispecific, as well as fragments, variants or derivatives thereof. Antibody fragments include those portions of the antibody which bind to an epitope on the IL-17 receptor like polypeptide. Examples of such fragments include Fab and F(ab') fragments generated by enzymatic cleavage of full-length antibodies. Other binding fragments include those generated by recombinant DNA techniques, such as



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the expression of recombinant plasmids containing nucleic acid sequences encoding antibody variable regions.

Polyclonal antibodies directed toward an IL-17 receptor like polypeptide generally are produced in animals (e.g., rabbits or mice) by means of multiple subcutaneous or intraperitoneal injections of IL-17 receptor like polypeptide and an adjuvant. It may be useful to conjugate an IL-17 receptor like polypeptide, including a variant, fragment, or derivative, to a carrier protein that is immunogenic in the species to be immunized, such as keyhole limpet heocyanin, serum, albumin, bovine thyroglobulin, or soybean trypsin inhibitor. Also, aggregating agents such as alum are used to enhance the immune response. After immunization, the animals are bled and the serum is assayed for anti-IL-17 receptor like polypeptide antibody titer.

Monoclonal antibodies directed toward an IL-17 receptor like polypeptide are produced using any method which provides for the production of antibody molecules by continuous cell lines in culture. Examples of suitable methods for preparing monoclonal antibodies include the hybridoma methods of Kohler et al., *Nature*, 256:495-497 (1975) and the human B-cell hybridoma method, Kozbor, J. *Immunol.*, 133:3001 (1984); Brodeur et al., *Monoclonal Antibody Production Techniques and Applications*, pp. 51-63 (Marcel Dekker, Inc., New York, 1987). Also provided by the invention are hybridoma cell lines which produce monoclonal antibodies reactive with IL-17 receptor like polypeptides.

Monoclonal antibodies of the invention may be modified for use as therapeutics. One embodiment is a "chimeric" antibody in which a portion of the heavy and/or light chain is identical with or homologous to a corresponding sequence in antibodies derived from a particular species or belonging to a particular antibody class or subclass, while the remainder of the chain(s) is identical with or homologous to a

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corresponding sequence in antibodies derived from another species or belonging to another antibody class or subclass. Also included are fragments of such antibodies, so long as they exhibit the desired biological activity. See, U.S. Patent No. 4,816,567; Morrison et al., *Proc. Natl. Acad. Sci.*, 81:6851-6855 (1985).

In another embodiment, a monoclonal antibody of the invention is a "humanized" antibody. Methods for humanizing non-human antibodies are well known in the art. See U.S. Patent Nos. 5,585,089, and 5,693,762. Generally, a humanized antibody has one or more amino acid residues introduced into it from a source which is non-human. Humanization can be performed, for example, using methods described in the art (Jones et al., *Nature* 321:522-525 (1986); Riechmann et al., *Nature*, 332:323-327 (1988); Verhoeyen et al., *Science* 239:1534-1536 (1988)), by substituting at least a portion of a rodent complementarity-determining region (CDR) for the corresponding regions of a human antibody.

Also encompassed by the invention are human antibodies which bind IL-17 receptor like polypeptides. Using transgenic animals (e.g., mice) that are capable of producing a repertoire of human antibodies in the absence of endogenous immunoglobulin production such antibodies are produced by immunization with an IL-17 receptor like antigen (i.e., having at least 6 contiguous amino acids), optionally conjugated to a carrier. See, for example, Jakobovits et al., *Proc. Natl. Acad. Sci.*, 90:2551-2555 (1993); Jakobovits et al., *Nature* 362:255-258 (1993); Bruggermann et al., *Year in Immuno.*, 7:33 (1993). In one method, such transgenic animals are produced by incapacitating the endogenous loci encoding the heavy and light immunoglobulin chains therein, and inserting loci encoding human heavy and light chain proteins into the genome thereof. Partially modified animals, that is those having less than the full complement of modifications, are then

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cross-bred to obtain an animal having all of the desired immune system modifications. When administered an immunogen, these transgenic animals produce antibodies with human variable regions, including human (rather than e.g., murine) amino acid sequences, including variable regions, including human which are immunospecific for these antigens. See PCT application nos. PCT/US96/05928 and PCT/US93/06926. Additional methods are described in U.S. Patent No. 5,545,807, PCT application nos. PCT/US91/245, PCT/GB89/01207, and in EP 546073B1 and EP 546073A1. Human antibodies may also be produced by the expression of recombinant DNA in host cells or by expression in hybridoma cells as described herein.

In an alternative embodiment, human antibodies can be produced from phage-display libraries (Hoogenboom et al., *J. Mol. Biol.* 227:381 (1991); Marks et al., *J. Mol. Biol.* 222:581 (1991). These processes mimic immune selection through the display of antibody repertoires on the surface of filamentous bacteriophage, and subsequent selection of phage by their binding to an antigen of choice. One such technique is described in PCT Application no. PCT/US98/17364, which describes the isolation of high affinity and functional agonistic antibodies for MPL- and msk- receptors using such an approach.

Chimeric, CDR grafted, and humanized antibodies are typically produced by recombinant methods. Nucleic acids encoding the antibodies are introduced into host cells and expressed using materials and procedures described herein. In a preferred embodiment, the antibodies are produced in mammalian host cells, such as CHO cells. Monoclonal (e.g., human) antibodies may be produced by the expression of recombinant DNA in host cells or by expression in hybridoma cells as described herein.

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The anti-IL-17 receptor like antibodies of the invention may be employed in any known assay method, such as competitive binding assays, direct and indirect sandwich assays, and immunoprecipitation assays (Sola, Monoclonal Antibodies: A  
5 Manual of Techniques, pp. 147-158 (CRC Press, Inc., 1987)) for the detection and quantitation of IL-17 receptor like polypeptides. The antibodies will bind IL-17 receptor like polypeptides with an affinity which is appropriate for the assay method being employed.

10 For diagnostic applications, in certain embodiments, anti-IL-17 receptor like antibodies may be labeled with a detectable moiety. The detectable moiety can be any one which is capable of producing, either directly or indirectly, a detectable signal. For example, the detectable moiety may be  
15 a radioisotope, such as  $^3\text{H}$ ,  $^{14}\text{C}$ ,  $^{32}\text{P}$ ,  $^{35}\text{S}$ , or  $^{125}\text{I}$ , a fluorescent or chemiluminescent compound, such as fluorescein isothiocyanate, rhodamine, or luciferin; or an enzyme, such as alkaline phosphatase,  $\beta$ -galactosidase, or horseradish peroxidase (Bayer et al., Meth. Enz., 184:138-163 (1990)).

20 Competitive binding assays rely on the ability of a labeled standard (e.g., an IL-17 receptor like polypeptide, or an immunologically reactive portion thereof) to compete with the test sample analyte (an IL-17 receptor like polypeptide) for binding with a limited amount of anti IL-17 receptor like  
25 antibody. The amount of an IL-17 receptor like polypeptide in the test sample is inversely proportional to the amount of standard that becomes bound to the antibodies. To facilitate determining the amount of standard that becomes bound, the antibodies typically are insolubilized before or after the  
30 competition, so that the standard and analyte that are bound to the antibodies may conveniently be separated from the standard and analyte which remain unbound.

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Sandwich assays typically involve the use of two antibodies, each capable of binding to a different immunogenic portion, or epitope, of the protein to be detected and/or quantitated. In a sandwich assay, the test sample analyte is typically bound by a first antibody which is immobilized on a solid support, and thereafter a second antibody binds to the analyte, thus forming an insoluble three part complex. See, e.g., U.S. Patent No. 4,376,110. The second antibody may itself be labeled with a detectable moiety (direct sandwich assays) or may be measured using an anti-immunoglobulin antibody that is labeled with a detectable moiety (indirect sandwich assays). For example, one type of sandwich assay is an enzyme-linked immunosorbent assay (ELISA), in which case the detectable moiety is an enzyme.

The selective binding agents, including anti-IL-17 receptor like antibodies, also are useful for *in vivo* imaging. An antibody labeled with a detectable moiety may be administered to an animal, preferably into the bloodstream, and the presence and location of the labeled antibody in the host is assayed. The antibody may be labeled with any moiety that is detectable in an animal, whether by nuclear magnetic resonance, radiology, or other detection means known in the art.

The invention also relates to a kit comprising IL-17 receptor like selective binding agents (such as antibodies) and other reagents useful for detecting IL-17 receptor like polypeptide levels in biological samples. Such reagents may include a secondary activity, a detectable label, blocking serum, positive and negative control samples, and detection reagents

Selective binding agents of the invention, including antibodies, may be used as therapeutics. These therapeutic agents are generally agonists or antagonists, in that they

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either enhance or reduce, respectively, at least one of the biological activities of an IL-17 receptor like polypeptide. In one embodiment, antagonist antibodies of the invention are antibodies or binding fragments thereof which are capable of specifically binding to an IL-17 receptor like polypeptide and which are capable of inhibiting or eliminating the functional activity of an IL-17 receptor like polypeptide *in vivo* or *in vitro*. In preferred embodiments, the selective binding agent, e.g., an antagonist antibody, will inhibit the functional activity of an IL-17 receptor like polypeptide by at least about 50%, and preferably by at least about 80%. In another embodiment, the selective binding agent may be an anti-IL-17 receptor like polypeptide antibody that is capable of interacting with an IL-17 receptor like binding partner (a ligand or receptor) thereby inhibiting or eliminating IL-17 receptor like activity *in vitro* or *in vivo*. Selective binding agents, including agonist and antagonist anti-IL-17 receptor like antibodies, are identified by screening assays which are well known in the art.

The invention also relates to a kit comprising IL-17 receptor like selective binding agents (such as antibodies) and other reagents useful for detecting IL-17 receptor like polypeptide levels in biological samples. Such reagents may include, a detectable label, blocking serum, positive and negative control samples, and detection reagents.

IL-17 receptor like polypeptides can be used to clone IL-17 receptor like ligand(s) using an "expression cloning" strategy. Radiolabeled (125-Iodine) IL-17 receptor like polypeptide or "affinity/activity-tagged" IL-17 receptor like polypeptide (such as an Fc fusion or an alkaline phosphatase fusion) can be used in binding assays to identify a cell type or cell line or tissue that expresses IL-17 receptor like ligand(s). RNA isolated from such cells or tissues can then be converted to cDNA, cloned into a mammalian expression

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vector, and transfected into mammalian cells (for example, COS, or 293) to create an expression library. Radiolabeled or tagged IL-17 receptor like polypeptide can then be used as an affinity reagent to identify and isolate the subset of cells in this library expressing IL-17 receptor like ligand(s). DNA is then isolated from these cells and transfected into mammalian cells to create a secondary expression library in which the fraction of cells expressing IL-17 receptor like ligand(s) would be many-fold higher than in the original library. This enrichment process can be repeated iteratively until a single recombinant clone containing an IL-17 receptor like ligand is isolated. Isolation of IL-17 receptor like ligand(s) is useful for identifying or developing novel agonists and antagonists of the IL-17 receptor like signaling pathway. Such agonists and antagonists include IL-17 receptor like ligand(s), anti-IL-17 receptor like ligand antibodies, small molecules or antisense oligonucleotides.

#### Assaying for Other Modulators of IL-17 Receptor Like Polypeptide Activity

In some situations, it may be desirable to identify molecules that are modulators, i.e., agonists or antagonists, of the activity of IL-17 receptor like polypeptide. Natural or synthetic molecules that modulate IL-17 receptor like polypeptide may be identified using one or more screening assays, such as those described herein. Such molecules may be administered either in an *ex vivo* manner, or in an *in vivo* manner by injection, or by oral delivery, implantation device, or the like.

"Test molecule(s)" refers to the molecule(s) that is/are under evaluation for the ability to modulate (i.e., increase or decrease) the activity of an IL-17 receptor like polypeptide. Most commonly, a test molecule will interact directly with an IL-17 receptor like polypeptide. However, it

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is also contemplated that a test molecule may also modulate IL-17 receptor like polypeptide activity indirectly, such as by affecting IL-17 receptor like gene expression, or by binding to an IL-17 receptor like binding partner (e.g., receptor or ligand). In one embodiment, a test molecule will bind to an IL-17 receptor like polypeptide with an affinity constant of at least about  $10^{-6}$  M, preferably about  $10^{-8}$  M, more preferably about  $10^{-9}$  M, and even more preferably about  $10^{-10}$  M.

10 Methods for identifying compounds which interact with IL-17 receptor like polypeptides are encompassed by the present invention. In certain embodiments, an IL-17 receptor like polypeptide is incubated with a test molecule under conditions which permit the interaction of the test molecule with an IL-17 receptor like polypeptide, and the extent of the interaction can be measured. The test molecule(s) can be screened in a substantially purified form or in a crude mixture. The test molecules can be nucleic acids, proteins, peptides, carbohydrates, lipids, organic and inorganic compounds.

In certain embodiments, an IL-17 receptor like polypeptide agonist or antagonist may be a protein, peptide, carbohydrate, lipid, or small molecular weight molecule which interacts with IL-17 receptor like polypeptide, or ligand thereof, to regulate its activity. Molecules which regulate IL-17 receptor like polypeptide expression include nucleic acids which are complementary to nucleic acids encoding an IL-17 receptor like polypeptide, or are complementary to nucleic acids sequences which direct or control the expression of IL-17 receptor like polypeptide, and which act as anti-sense regulators of expression.

Once a set of test molecules has been identified as interacting with an IL-17 receptor like polypeptide, the



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molecules may be further evaluated for their ability to increase or decrease IL-17 receptor like polypeptide activity. The measurement of the interaction of test molecules with IL-17 receptor like polypeptides may be carried out in several formats, including cell-based binding assays, membrane binding assays, solution-phase assays and immunoassays. In general, test molecules are incubated with an IL-17 receptor like polypeptide for a specified period of time, and IL-17 receptor like polypeptide activity is determined by one or more assays for measuring biological activity.

The interaction of test molecules with IL-17 receptor like polypeptides may also be assayed directly using polyclonal or monoclonal antibodies in an immunoassay. Alternatively, modified forms of IL-17 receptor like polypeptides containing epitope tags as described herein may be used in immunoassays.

In certain embodiments, a IL-17 receptor like polypeptide agonist or antagonist may be a protein, peptide, carbohydrate, lipid, or small molecular weight molecule which interacts with IL-17 receptor like polypeptide to regulate its activity. Potential protein antagonists of IL-17 receptor like polypeptide include antibodies which interact with active regions of the polypeptide and inhibit or eliminate at least one activity of IL-17 receptor like molecules. Molecules which regulate IL-17 receptor like polypeptide expression include nucleic acids which are complementary to nucleic acids encoding a IL-17 receptor like polypeptide, or are complementary to nucleic acids sequences which direct or control the expression of IL-17 receptor like polypeptide, and which act as anti-sense regulators of expression.

In the event that IL-17 receptor like polypeptides display biological activity through an interaction with a binding partner (e.g., selective binding agent or a ligand), a

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variety of *in vitro* assays may be used to measure the binding of an IL-17 receptor like polypeptide to the corresponding binding partner (such as a selective binding agent or ligand). These assays may be used to screen test molecules for their ability to increase or decrease the rate and/or the extent of binding of an IL-17 receptor like polypeptide to its binding partner. In one assay, an IL-17 receptor like polypeptide is immobilized in the wells of a microtiter plate. Radiolabeled IL-17 receptor like binding partner (for example, iodinated IL-17 receptor like binding partner) and the test molecule(s) can then be added either one at a time (in either order) or simultaneously to the wells. After incubation, the wells can be washed and counted, using a scintillation counter, for radioactivity to determine the extent to which the binding partner bound to IL-17 receptor like polypeptide. Typically, the molecules will be tested over a range of concentrations, and a series of control wells lacking one or more elements of the test assays can be used for accuracy in the evaluation of the results. An alternative to this method involves reversing the "positions" of the proteins, i.e., immobilizing IL-17 receptor like binding partner to the microtiter plate wells, incubating with the test molecule and radiolabeled IL-17 receptor like polypeptide, and determining the extent of IL-17 receptor like polypeptide binding. See, for example, chapter 18, *Current Protocols in Molecular Biology*, Ausubel et al., eds., John Wiley & Sons, New York, NY (1995).

As an alternative to radiolabelling, an IL-17 receptor like polypeptide or its binding partner may be conjugated to biotin and the presence of biotinylated protein can then be detected using streptavidin linked to an enzyme, such as horseradish peroxidase (HRP) or alkaline phosphatase (AP), that can be detected colorometrically, or by fluorescent tagging of streptavidin. An antibody directed to an IL-17 receptor like polypeptide or to an IL-17 receptor like binding partner and conjugated to biotin may also be used and can be

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detected after incubation with enzyme-linked streptavidin linked to AP or HRP.

An IL-17 receptor like polypeptide or an IL-17 receptor like binding partner can also be immobilized by attachment to  
5 agarose beads, acrylic beads or other types of such inert solid phase substrates. The substrate-protein complex can be placed in a solution containing the complementary protein and the test compound. After incubation, the beads can be precipitated by centrifugation, and the amount of binding  
10 between an IL-17 receptor like polypeptide and its binding partner can be assessed using the methods described herein. Alternatively, the substrate-protein complex can be immobilized in a column, and the test molecule and complementary protein are passed through the column. The  
15 formation of a complex between an IL-17 receptor like polypeptide and its binding partner can then be assessed using any of the techniques set forth herein, i.e., radiolabelling, antibody binding, or the like.

Another *in vitro* assay that is useful for identifying a  
20 test molecule which increases or decreases the formation of a complex between an IL-17 receptor like binding protein and an IL-17 receptor like binding partner is a surface plasmon resonance detector system such as the BIAcore assay system (Pharmacia, Piscataway, NJ). The BIAcore system may be  
25 carried out using the manufacturer's protocol. This assay essentially involves the covalent binding of either IL-17 receptor like polypeptide or an IL-17 receptor like binding partner to a dextran-coated sensor chip which is located in a detector. The test compound and the other complementary  
30 protein can then be injected, either simultaneously or sequentially, into the chamber containing the sensor chip. The amount of complementary protein that binds can be assessed based on the change in molecular mass which is physically associated with the dextran-coated side of the sensor chip;

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the change in molecular mass can be measured by the detector system.

In some cases, it may be desirable to evaluate two or more test compounds together for their ability to increase or decrease the formation of a complex between an IL-17 receptor like polypeptide and an IL-17 receptor like binding partner. In these cases, the assays set forth herein can be readily modified by adding such additional test compound(s) either simultaneous with, or subsequent to, the first test compound. The remainder of the steps in the assay are as set forth herein.

*In vitro* assays such as those described herein may be used advantageously to screen large numbers of compounds for effects on complex formation by IL-17 receptor like polypeptide and IL-17 receptor like binding partner. The assays may be automated to screen compounds generated in phage display, synthetic peptide, and chemical synthesis libraries.

Compounds which increase or decrease the formation of a complex between an IL-17 receptor like polypeptide and an IL-17 receptor like binding partner may also be screened in cell culture using cells and cell lines expressing either IL-17 receptor like polypeptide or IL-17 receptor like binding partner. Cells and cell lines may be obtained from any mammal, but preferably will be from human or other primate, canine, or rodent sources. The binding of an IL-17 receptor like polypeptide to cells expressing IL-17 receptor like binding partner at the surface is evaluated in the presence or absence of test molecules, and the extent of binding may be determined by, for example, flow cytometry using a biotinylated antibody to an IL-17 receptor like binding partner. Cell culture assays can be used advantageously to further evaluate compounds that score positive in protein binding assays described herein.

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Cell cultures can also be used to screen the impact of a drug candidate. For example, drug candidates may decrease or increase the expression of the IL-17 receptor like gene. In certain embodiments, the amount of IL-17 receptor like polypeptide or a fragment that is produced may be measured after exposure of the cell culture to the drug candidate. In certain embodiments, one may detect the actual impact of the drug candidate on the cell culture. For example, the overexpression of a particular gene may have a particular impact on the cell culture. In such cases, one may test a drug candidate's ability to increase or decrease the expression of the gene or its ability to prevent or inhibit a particular impact on the cell culture. In other examples, the production of a particular metabolic product such as a fragment of a polypeptide, may result in, or be associated with, a disease or pathological condition. In such cases, one may test a drug candidate's ability to decrease the production of such a metabolic product in a cell culture.

A yeast two hybrid system (Chien et al., Proc. Natl. Acad. Sci. USA, 88: 9578-9583, 1991) can be used to identify novel polypeptides that bind to a yeast-two hybrid bait construct can be generated in a vector (such as the pAS2-1 form Clontech) which encodes a yeast-two hybrid domain fused to the IL-17 receptor like polynucleotide. This bait construct may be used to screen human cDNA libraries wherein the cDNA library sequences are fused to GAL4 activation domains. Positive interactions will result in the activation of a reporter gene such as  $\beta$ -gal. Positive clones emerging from the screening may be characterized further to identify interacting proteins.

### Internalizing Proteins

The tat protein sequence (from HIV) can be used to internalize proteins into a cell. See e.g., Falwell et al.,

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*Proc. Natl. Acad. Sci.*, 91:664-668 (1994). For example, an 11 amino acid sequence (YGRKKRRQRRR; SEQ ID NO: 16) of the HIV tat protein (termed the "protein transduction domain", or TAT PDT) has been described as mediating delivery across the cytoplasmic membrane and the nuclear membrane of a cell. See Schwarze et al., *Science*, 285:1569-1572 (1999); and Nagahara et al., *Nature Medicine*, 4:1449-1452 (1998). In these procedures, FITC-constructs (FITC-GGGGYGRKKRRQRRR; SEQ ID NO: 17) are prepared which bind to cells as observed by fluorescence-activated cell sorting (FACS) analysis, and these constructs penetrate tissues after i.p. administration. Next, tat-bgal fusion proteins are constructed. Cells treated with this construct demonstrated  $\beta$ -gal activity. Following injection, a number of tissues, including liver, kidney, lung, heart, and brain tissue have been found to demonstrate expression using these procedures. It is believed that these constructions underwent some degree of unfolding in order to enter the cell; as such, refolding may be required after entering the cell.

20

It will thus be appreciated that the tat protein sequence may be used to internalize a desired protein or polypeptide into a cell. For example, using the tat protein sequence, an IL-17 receptor like antagonist (such as an anti-IL-17 receptor like selective binding agent, small molecule, soluble receptor, or antisense oligonucleotide) can be administered intracellularly to inhibit the activity of an IL-17 receptor like molecule. As used herein, the term "IL-17 receptor like molecule" refers to both IL-17 receptor like nucleic acid molecules and IL-17 receptor like polypeptides as defined herein. Where desired, the IL-17 receptor like protein itself may also be internally administered to a cell using these procedures. See also, Strauss, E., "Introducing Proteins Into the Body's Cells", *Science*, 285:1466-1467 (1999).

Cell Source Identification Using IL-17 Receptor Like Polypeptides

In accordance with certain embodiments of the invention, it may be useful to be able to determine the source of a certain cell type associated with an IL-17 receptor like polypeptide. For example, it may be useful to determine the origin of a disease or pathological condition as an aid in selecting an appropriate therapy.

Therapeutic Uses

A non-exclusive list of acute and chronic diseases which can be treated, diagnosed, ameliorated, or prevented with the IL-17 receptor like nucleic acids, polypeptides, and agonists and antagonists of the invention include:

- The diagnosis and/or treatment of diseases involving immune system dysfunction. Examples of such diseases include, but are not limited to, rheumatoid arthritis, psoriatic arthritis, inflammatory arthritis, osteoarthritis, inflammatory joint disease, autoimmune disease including autoimmune vasculitis, multiple sclerosis, lupus, diabetes (e.g., insulin diabetes), inflammatory bowel disease, transplant rejection, graft vs. host disease, and inflammatory conditions resulting from strain, sprain, cartilage damage, trauma, orthopedic surgery, infection or other disease processes. Other diseases influenced by the dysfunction of the immune system are encompassed within the scope of the invention, including but not limited to, allergies. The IL-17 receptor like nucleic acids, polypeptides, and agonists and antagonists of the invention can also be used to inhibit T cell proliferation, to inhibit T cell activation, and/or to inhibit B cell proliferation and/or immunoglobulin secretion.

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- The diagnosis and/or treatment of diseases involving infection. Examples of such diseases include, but are not limited to, leprosy, viral infections such as hepatitis or HIV, bacterial infection such as *clostridium* associated illnesses, including *clostridium*-associated diarrhea, pulmonary tuberculosis, acute febrile illness from bacteria such as or virus, fever, acute phase response of the liver, septicemia, septic shock. Other diseases involving infection are encompassed within the scope of the invention.
- The diagnosis and/or treatment of diseases involving weight disorders. Examples of such diseases include, but are not limited to obesity, anorexia, cachexia, including AIDS-induced cachexia, myopathies (e.g., muscle protein metabolism, such as in sepsis), and hypoglycemia. Other diseases involving weight disorders are encompassed within the scope of the invention.
- The diagnosis and/or treatment of diseases involving neuronal dysfunction. Examples of such diseases include, but are not limited to Alzheimer's, Parkinson's disease, neurotoxicity (e.g., as induced by HIV), ALS, brain injury, stress, depression, nociception and other pain (including cancer-related pain), hyperalgesia, epilepsy, learning impairment and memory disorders, sleep disturbance, and peripheral and central neuropathies. Other neurological disorders are encompassed within the scope of the invention.
- The diagnosis and/or treatment of diseases involving the lung. Examples of such diseases include, but are not limited to, acute or chronic lung injury including interstitial lung disease, acute respiratory disease syndrome, pulmonary hypertension, emphysema, cystic fibrosis, pulmonary fibrosis, and asthma. Other diseases of the lung are encompassed within the scope of the



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invention.

- 5     • The diagnosis and/or treatment of diseases involving the skin. Examples of such diseases include, but are not limited to, psoriasis, eczema, and wound healing. Other diseases of the skin are encompassed within the scope of the invention.
- 10    • The diagnosis and/or treatment of diseases involving the kidney. Examples of such diseases include, but are not limited to, acute and chronic glomerulonephritis. Other diseases of the kidney are encompassed within the scope of the invention.
- 15    • The diagnosis and/or treatment of diseases involving the bone. Examples of such diseases include, but are not limited to, ~~osteoporosis, osteopetrosis, osteogenesis~~ imperfecta, Paget's disease, periodontal disease, temporal mandibular joint disease, and hypercalcemia. Other diseases of the bone are encompassed within the scope of the invention.
- 20    • The diagnosis and/or treatment of diseases involving the vascular system. Examples of such diseases include, but are not limited to hemorrhage or stroke, hemorrhagic shock, ischemia, including cardiac ischemia and cerebral ischemia (e.g., brain injury as a result of trauma, 25    epilepsy, hemorrhage or stroke, each of which may lead to neurodegeneration), atherosclerosis, congestive heart failure; restenosis, reperfusion injury, and angiogenesis. Other diseases of the vascular system are encompassed within the scope of the invention.
- 30    • The diagnosis and/or treatment of tumor cells. Examples of such diseases include, but are not limited to, lymphomas, bone sarcoma, chronic and acute myelogenous leukemia (AML and CML) and other leukemias, multiple

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myeloma, lung, breast cancer, tumor metastasis, and side effects from radiation therapy. Other diseases involving tumor cells are encompassed within the scope of the invention.

- 5       • The diagnosis and/or treatment of reproductive disorders. Examples of such diseases include, but are not limited to, infertility, miscarriage, pre-term labor and delivery, and endometriosis. Other diseases involving the reproductive system are encompassed within the scope of the invention.
- 10
- The diagnosis and/or treatment of eye disorders. Examples of such diseases include, but are not limited to, inflammatory eye disease, as may be associated with, for example, corneal transplant; retinal degeneration, blindness, macular degeneration, glaucoma, uveitis, and retinal neuropathy. Other diseases of the eye are encompassed within the scope of the invention.
- 15

20       Other diseases which are treatable using agents within the scope of the invention include acute pancreatitis, chronic fatigue syndrome, fibromyalgia, and Kawasaki's disease (MLNS).

      Other diseases associated with undesirable levels of one or more of IL-1, IL-1ra, the ligand of the present IL-17 receptor like polypeptide, and/or the present IL-17 receptor like polypeptide itself are encompassed within the scope of the invention. Undesirable levels include excessive and/or sub-normal levels of IL-1, IL-1ra, the ligand of the present IL-17 receptor like polypeptide, and/or the IL-17 receptor like polypeptides described herein.

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      IL-1 inhibitors include any protein capable of specifically preventing activation of cellular receptors to

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IL-1, which may result from any number of mechanisms. Such mechanisms include downregulating IL-1 production, binding free IL-1, interfering with IL-1 binding to its receptor, interfering with formation of the IL-1 receptor complex (i.e., association of IL-1 receptor with IL-1 receptor accessory protein), or interfering with modulation of IL-1 signaling after binding to its receptor. Classes of interleukin-1 inhibitors include:

- 10 • Interleukin-1 receptor antagonists such as IL-1ra, as described herein;
- Anti-IL-1 receptor monoclonal antibodies (e.g., EP 623674);
- 15 • IL-1 binding proteins such as soluble IL-1 receptors (e.g., U.S. Pat. No. 5,492,888, U.S. Pat. No. 5,488,032, and U.S. Pat. No. 5,464,937, U.S. Pat. No. 5,319,071, and U.S. Pat. No. 5,180,812;
- Anti-IL-1 monoclonal antibodies (e.g., WO 9501997, WO 9402627, WO 9006371, U.S. Pat. No. 4,935,343, EP 364778, EP 267611 and EP 220063;
- 20 • IL-1 receptor accessory proteins and antibodies thereto (e.g., WO 96/23067);
- Inhibitors of interleukin-1 $\beta$  converting enzyme (ICE) or caspase I, which can be used to inhibit IL-1 beta production and secretion;
- 25 • Interleukin-1 $\beta$  protease inhibitors;
- Other compounds and proteins which block *in vivo* synthesis or extracellular release of IL-1.

Exemplary IL-1 inhibitors are disclosed in the following references:

- 30 US Pat. Nos. 5747444; 5359032; 5608035; 5843905; 5359032; 5866576; 5869660; 5869315; 5872095; 5955480;

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International (WO) patent applications 98/21957,  
96/09323, 91/17184, 96/40907, 98/32733, 98/42325, 98/44940,  
98/47892, 98/56377, 99/03837, 99/06426, 99/06042, 91/17249,  
98/32733, 98/17661, 97/08174, 95/34326, 99/36426, and  
5 99/36415;

European (EP) patent applications 534978 and 894795; and  
French patent application FR 2762514;

Interleukin-1 receptor antagonist (IL-1ra) is a human  
protein that acts as a natural inhibitor of interleukin-1.  
10 Preferred receptor antagonists (including IL-1ra and variants  
and derivatives thereof), as well as methods of making and  
using thereof, are described in U.S. Patent No. 5,075,222; WO  
91/08285; WO 91/17184; AU 9173636; WO 92/16221; WO93/21946; WO  
94/06457; WO 94/21275; FR 2706772; WO 94/21235; DE 4219626, WO  
15 94/20517; WO 96/22793; WO 97/28828; and WO 99/36541. The  
proteins include glycosylated as well as non-glycosylated IL-1  
receptor antagonists.

Specifically, three exemplary forms of IL-1ra and  
variants thereof are disclosed and described in the 5,075,222  
20 patent. The first of these, called "IL-1i" in the 5,075,222  
patent, is characterized as a 22-23 kD molecule on SDS-PAGE  
with an approximate isoelectric point of 4.8, eluting from a  
MonoQ FPLC column at around 52 mM NaCl in Tris buffer, pH 7.6.  
The second, IL-1ra $\beta$ , is characterized as a 22-23 kD protein,  
25 eluting from a MonoQ column at 48 mM NaCl. Both IL-1ra $\alpha$  and  
IL-1ra $\beta$  are glycosylated. The third, IL-1rax, is  
characterized as a 20 kD protein, eluting from a MonoQ column  
at 48 mM NaCl, and is non-glycosylated. U.S. patent No.  
5,075,222 also discloses methods for isolating the genes  
30 responsible for coding the inhibitors, cloning the gene in  
suitable vectors and cell types, and expressing the gene to  
produce the inhibitors.

Those skilled in the art will recognize that many  
combinations of deletions, insertions, and substitutions

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(individually or collectively "variant(s)" herein) can be made within the amino acid sequences of IL-1ra, provided that the resulting molecule is biologically active (e.g., possesses the ability to affect one or more of the diseases and disorders such as those recited herein.)

As contemplated by the present invention, an agonist or antagonist of the IL-17 receptor like polypeptide (including, but not limited to, anti-IL-17 receptor like selective binding agents [such as antibodies], ligands to the IL-17 RECEPTOR like receptor, soluble IL-17 receptor like polypeptides, small molecules, and antisense oligonucleotides, or an IL-17 receptor like polypeptide itself) may be administered as an adjunct to other therapy and also with other pharmaceutical compositions suitable for the indication being treated. An agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself and any of one or more additional therapies or pharmaceutical formulations may be administered separately, sequentially, or simultaneously.

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In a specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pre-treatment, post-treatment, or concurrent treatment) with any of one or more TNF inhibitors for the treatment or prevention of the diseases and disorders recited herein.

Such TNF inhibitors include compounds and proteins which block in vivo synthesis or extracellular release of TNF. In a specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pre-treatment, post-treatment, or concurrent treatment) with any of one or more of the following TNF

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inhibitors: TNF binding proteins (soluble TNF receptor type-I and soluble TNF receptor type-II ("sTNFRs"), as defined herein), anti-TNF antibodies, granulocyte colony stimulating factor; thalidomide; BN 50730; tenidap; E 5531; tiapafant PCA  
5 4248; nimesulide; panavir; rolipram; RP 73401; peptide T; MDL 201,449A; (1R,3S)-Cis-1-[9-(2,6-diaminopurinyl)]-3-hydroxy-4-cyclopentene hydrochloride; (1R,3R)-trans-1-(9-(2,6-diamino)purine]-3-acetoxycyclopentane; (1R,3R)-trans-1-[9-adenyl)-3-azidocyclopentane hydrochloride and (1R,3R)-trans-1-  
10 (6-hydroxy-purin-9-yl)-3-azidocyclopentane. TNF binding proteins are disclosed in the art (EP 308 378, EP 422 339, GB 2 218 101, EP 393 438, WO 90/13575, EP 398 327, EP 412 486, WO 91/03553, EP 418 014, JP 127,800/1991, EP 433 900, U.S. Patent No. 5,136,021, GB 2 246 569, EP 464 533, WO 92/01002,  
15 WO 92/13095, WO 92/16221, EP 512 528, EP 526 905, WO 93/07863, EP 568 928, WO 93/21946, WO 93/19777, EP 417 563, WO 94/06476, and PCT International Application No. PCT/US97/12244).

For example, EP 393 438 and EP 422 339 teach the amino acid and nucleic acid sequences of a soluble TNF receptor type  
20 I (also known as "sTNFR-I" or "30kDa TNF inhibitor") and a soluble TNF receptor type II (also known as "sTNFR-II" or "40kDa TNF inhibitor"), collectively termed "sTNFRs", as well as modified forms thereof (e.g., fragments, functional derivatives and variants). EP 393 438 and EP 422 339 also  
25 disclose methods for isolating the genes responsible for coding the inhibitors, cloning the gene in suitable vectors and cell types and expressing the gene to produce the inhibitors. Additionally, polyvalent forms (i.e., molecules comprising more than one active moiety) of sTNFR-I and sTNFR-  
30 II have also been disclosed. In one embodiment, the polyvalent form may be constructed by chemically coupling at least one TNF inhibitor and another moiety with any clinically acceptable linker, for example polyethylene glycol (WO 92/16221 and WO 95/34326), by a peptide linker (Neve et  
35 al. (1996), *Cytokine*, 8(5):365-370, by chemically coupling to

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biotin and then binding to avidin (WO 91/03553) and, finally, by combining chimeric antibody molecules (U.S. Patent 5,116,964, WO 89/09622, WO 91/16437 and EP 315062).

Anti-TNF antibodies include MAK 195F Fab antibody (Holler  
5 et al. (1993), 1st International Symposium on Cytokines in  
Bone Marrow Transplantation, 147); CDP 571 anti-TNF monoclonal  
antibody (Rankin et al. (1995), *British Journal of*  
*Rheumatology*, 34:334-342); BAY X 1351 murine anti-tumor  
necrosis factor monoclonal antibody (Kieft et al. (1995),  
10 7th European Congress of Clinical Microbiology and Infectious  
Diseases, page 9); CentTNF cA2 anti-TNF monoclonal antibody  
(Elliott et al. (1994), *Lancet*, 344:1125-1127 and Elliott et  
al. (1994), *Lancet*, 344:1105-1110).

In a specific embodiment, the present invention is  
15 directed to the use of agonist or antagonist of the IL-17  
receptor like polypeptide, and/or an IL-17 receptor like  
polypeptide itself in combination (pretreatment, post-  
treatment, or concurrent treatment) with secreted or soluble  
human fas antigen or recombinant versions thereof (WO 96/20206  
20 and Mountz et al., *J. Immunology*, 155:4829-4837; and EP 510  
691. WO 96/20206 discloses secreted human fas antigen (native  
and recombinant, including an Ig fusion protein), methods for  
isolating the genes responsible for coding the soluble  
recombinant human fas antigen, methods for cloning the gene in  
25 suitable vectors and cell types, and methods for expressing  
the gene to produce the inhibitors. EP 510 691 describes DNAs  
coding for human fas antigen, including soluble fas antigen,  
vectors expressing for said DNAs and transformants transfected  
with the vector. When administered parenterally, doses of a  
30 secreted or soluble fas antigen fusion protein each are  
generally from about 1 micrograms/kg to about 100  
micrograms/kg.

Treatment of the diseases and disorders recited herein,  
including acute and chronic inflammation such as rheumatic

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diseases, commonly includes the use of first line drugs for control of pain and inflammation; these drugs are classified as non-steroidal, anti-inflammatory drugs (NSAIDs). Secondary treatments include corticosteroids, slow acting antirheumatic drugs (SAARDs), or disease modifying (DM) drugs. Information regarding the following compounds can be found in The Merck Manual of Diagnosis and Therapy, Sixteenth Edition, Merck, Sharp & Dohme Research Laboratories, Merck & Co., Rahway, NJ (1992) and in Pharmaprojects, PJB Publications Ltd.

10 In a specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself and any of one or more NSAIDs for the treatment of the diseases and disorders recited herein, including acute and chronic inflammation such as rheumatic diseases; and graft versus host disease. NSAIDs owe their anti-inflammatory action, at least in part, to the inhibition of prostaglandin synthesis (Goodman and Gilman in "The Pharmacological Basis of Therapeutics," MacMillan 7th Edition  
15 (1985)). NSAIDs can be characterized into at least nine groups: (1) salicylic acid derivatives; (2) propionic acid derivatives; (3) acetic acid derivatives; (4) fenamic acid derivatives; (5) carboxylic acid derivatives; (6) butyric acid derivatives; (7) oxicams; (8) pyrazoles and (9) pyrazolones.

25 In another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more salicylic acid derivatives, prodrug esters or pharmaceutically acceptable salts thereof. Such salicylic acid derivatives, prodrug esters and pharmaceutically acceptable salts thereof comprise: acetaminosalol, aloxiprin, aspirin, benorylate, bromosaligenin, calcium acetylsalicylate, choline magnesium  
30 trisalicylate, magnesium salicylate, choline salicylate,



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diflusal, etersalate, fendosal, gentisic acid, glycol salicylate, imidazole salicylate, lysine acetylsalicylate, mesalamine, morpholine salicylate, 1-naphthyl salicylate, olsalazine, parsalimide, phenyl acetylsalicylate, phenyl salicylate, salacetamide, salicylamide O-acetic acid, salsalate, sodium salicylate and sulfasalazine. Structurally related salicylic acid derivatives having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

10 In an additional specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or  
15 more propionic acid derivatives, prodrug esters or pharmaceutically acceptable salts thereof. The propionic acid derivatives, prodrug esters, and pharmaceutically acceptable salts thereof comprise: alminoprofen, benoxaprofen, bucloxic acid, carprofen, dexindoprofen, fenoprofen, flunoxaprofen, fluprofen, flurbiprofen, furocloprofen, ibuprofen, ibuprofen aluminum, ibuprofen, indoprofen, isoprofen, ketoprofen, loxoprofen, miroprofen, naproxen, naproxen sodium, oxaprozin, piketoprofen, pimeprofen, pirprofen, pranoprofen, protizinic acid, pyridoxiprofen, suprofen, tiaprofenic acid and  
20 tioxaprofen. Structurally related propionic acid derivatives having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

In yet another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-  
30 17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more acetic acid derivatives, prodrug esters or pharmaceutically acceptable salts thereof. The acetic acid derivatives,  
35 prodrug esters, and pharmaceutically acceptable salts thereof

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comprise: acetamin, alclofenac, amfenac, bufexamac, cinmetacin, clopirac, delmetacin, diclofenac potassium, diclofenac sodium, etodolac, felbinac, fenclofenac, fenclozic acid, fentiazac, furofenac, glucametacin, ibufenac, 5 indomethacin, isofezolac, isoxepac, lonazolac, metiazinic acid, oxametacin, oxpinac, pimetacin, proglumetacin, sulindac, talmetacin, tiaramide, tiopinac, tolmetin, tolmetin sodium, zidometacin and zomepirac. Structurally related acetic acid derivatives having similar analgesic and anti-inflammatory 10 properties are also intended to be encompassed by this group.

In another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post- 15 treatment, or concurrent treatment) with any of one or more fenamic acid derivatives, prodrug esters or pharmaceutically acceptable salts thereof. The fenamic acid derivatives, prodrug esters and pharmaceutically acceptable salts thereof comprise: enfenamic acid, etofenamate, flufenamic acid, 20 isonixin, meclofenamic acid, meclofenamate sodium, medofenamic acid, mefenamic acid, niflumic acid, talniflumate, terofenamate, tolfenamic acid and ufenamate. Structurally related fenamic acid derivatives having similar analgesic and anti-inflammatory properties are also intended to be 25 encompassed by this group.

In an additional specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, 30 post-treatment, or concurrent treatment) with any of one or more carboxylic acid derivatives, prodrug esters or pharmaceutically acceptable salts thereof. The carboxylic acid derivatives, prodrug esters, and pharmaceutically acceptable salts thereof which can be used comprise: clidanac, 35 diflunisal, flufenisal, inoridine, ketorolac and tinoridine.

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Structurally related carboxylic acid derivatives having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

In yet another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more butyric acid derivatives, prodrug esters or pharmaceutically acceptable salts thereof. The butyric acid derivatives, prodrug esters, and pharmaceutically acceptable salts thereof comprise: bumadizon, butibufen, fenbufen and xenbucin. Structurally related butyric acid derivatives having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

In another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more oxicams, prodrug esters, or pharmaceutically acceptable salts thereof. The oxicams, prodrug esters, and pharmaceutically acceptable salts thereof comprise: droxicam, enolicam, isoxicam, piroxicam, sudoxicam, tenoxicam and 4-hydroxyl-1,2-benzothiazine-1,1-dioxide-4-(N-phenyl)-carboxamide. Structurally related oxicams having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

In still another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more pyrazoles, prodrug esters, or pharmaceutically acceptable

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salts thereof. The pyrazoles, prodrug esters, and pharmaceutically acceptable salts thereof which may be used comprise: difenamizole and epirizole. Structurally related pyrazoles having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

In an additional specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment or, concurrent treatment) with any of one or more pyrazolones, prodrug esters, or pharmaceutically acceptable salts thereof. The pyrazolones, prodrug esters and pharmaceutically acceptable salts thereof which may be used comprise: apazone, azapropazone, benzpiperylon, feprazone, mofebutazone, morazone, oxyphenbutazone, phenylbutazone, pipebuzone, propylphenazone, ramifenazone, suxibuzone and thiazolinobutazone. Structurally related pyrazalones having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

In another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more of the following NSAIDs:  $\epsilon$ -acetamidocaproic acid, S-adenosylmethionine, 3-amino-4-hydroxybutyric acid, amixetrine, anitrazafen, antrafenine, bendazac, bendazac lysinate, benzydamine, beprozoin, broperamole, bucolome, bufezolac, ciproquazone, cloximate, dazidamine, deboxamet, detomidine, difenpiramide, difenpyramide, difisalamine, ditazol, emorfazone, fanetizole mesylate, fenflumizole, floctafenine, flumizole, flunixin, fluproquazone, fopirtoline, fosfosal, guaimesal, guaiazolene, isonixirn, lefetamine HCl,

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leflunomide, lofemizole, lotifazole, lysin clonixinate, meseclazone, nabumetone, nictindole, nimesulide, orgotein, orpanoxin, oxaceprol, oxapadol, paranyline, perisoxal, perisoxal citrate, pifoxime, piroxifen, pirazolac, pirfenidone, proquazone, proxazole, thielavin B, tiflamizole, timegadine, tolectin, tolpadol, tryptamid and those designated by company code number such as 480156S, AA861, AD1590, AFP802, AFP860, AI77B, AP504, AU8001, BPPC, BW540C, CHINOIN 127, CN100, EB382, EL508, F1044, FK-506, GV3658, ITF182, KCNTEI6090, KME4, LA2851, MR714, MR897, MY309, ONO3144, PR823, PV102, PV108, R830, RS2131, SCR152, SH440, SIR133, SPAS510, SQ27239, ST281, SY6001, TA60, TAI-901 (4-benzoyl-1-indancarboxylic acid), TVX2706, U60257, UR2301 and WY41770. Structurally related NSAIDs having similar analgesic and anti-inflammatory properties to the NSAIDs are also intended to be encompassed by this group.

In still another specific embodiment, the present invention is directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment or concurrent treatment) with any of one or more corticosteroids, prodrug esters or pharmaceutically acceptable salts thereof for the treatment of the diseases and disorders recited herein, including acute and chronic inflammation such as rheumatic diseases, graft versus host disease and multiple sclerosis. Corticosteroids, prodrug esters and pharmaceutically acceptable salts thereof include hydrocortisone and compounds which are derived from hydrocortisone, such as 21-acetoxypregnenolone, alclomerasone, algestone, amcinonide, beclomethasone, betamethasone, betamethasone valerate, budesonide, chloroprednisone, clobetasol, clobetasol propionate, clobetasone, clobetasone butyrate, clocortolone, clocprednol, corticosterone, cortisone, cortivazol, deflazacort, desonide, desoximetasone, dexamethasone, diflorasone, diflucortolone, difluprednate,

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enoxolone, fluazacort, flucoronide, flumethasone, flumethasone pivalate, flucinolone acetonide, flunisolide, fluocinonide, fluorocinolone acetonide, fluocortin butyl, fluocortolone, fluocortolone hexanoate, difluocortolone  
5 valerate, fluorometholone, fluperolone acetate, fluprednidene acetate, fluprednisolone, flurandenolide, formocortal, halcinonide, halometasone, halopredone acetate, hydrocortamate, hydrocortisone, hydrocortisone acetate, hydrocortisone butyrate, hydrocortisone phosphate,  
10 hydrocortisone 21-sodium succinate, hydrocortisone tebutate, mazipredone, medrysone, meprednisone, methylprednisolone, mometasone furoate, paramethasone, prednicarbate, prednisolone, prednisolone 21-diedryaminoacetate, prednisolone sodium phosphate, prednisolone sodium succinate, prednisolone  
15 sodium 21-m-sulfobenzoate, prednisolone sodium 21-stearoglycolate, prednisolone tebutate, prednisolone 21-trimethylacetate, prednisone, prednival, prednylidene, prednylidene 21-diethylaminoacetate, tixocortol, triamcinolone, triamcinolone acetonide, triamcinolone  
20 benetonide and triamcinolone hexacetonide. Structurally related corticosteroids having similar analgesic and anti-inflammatory properties are also intended to be encompassed by this group.

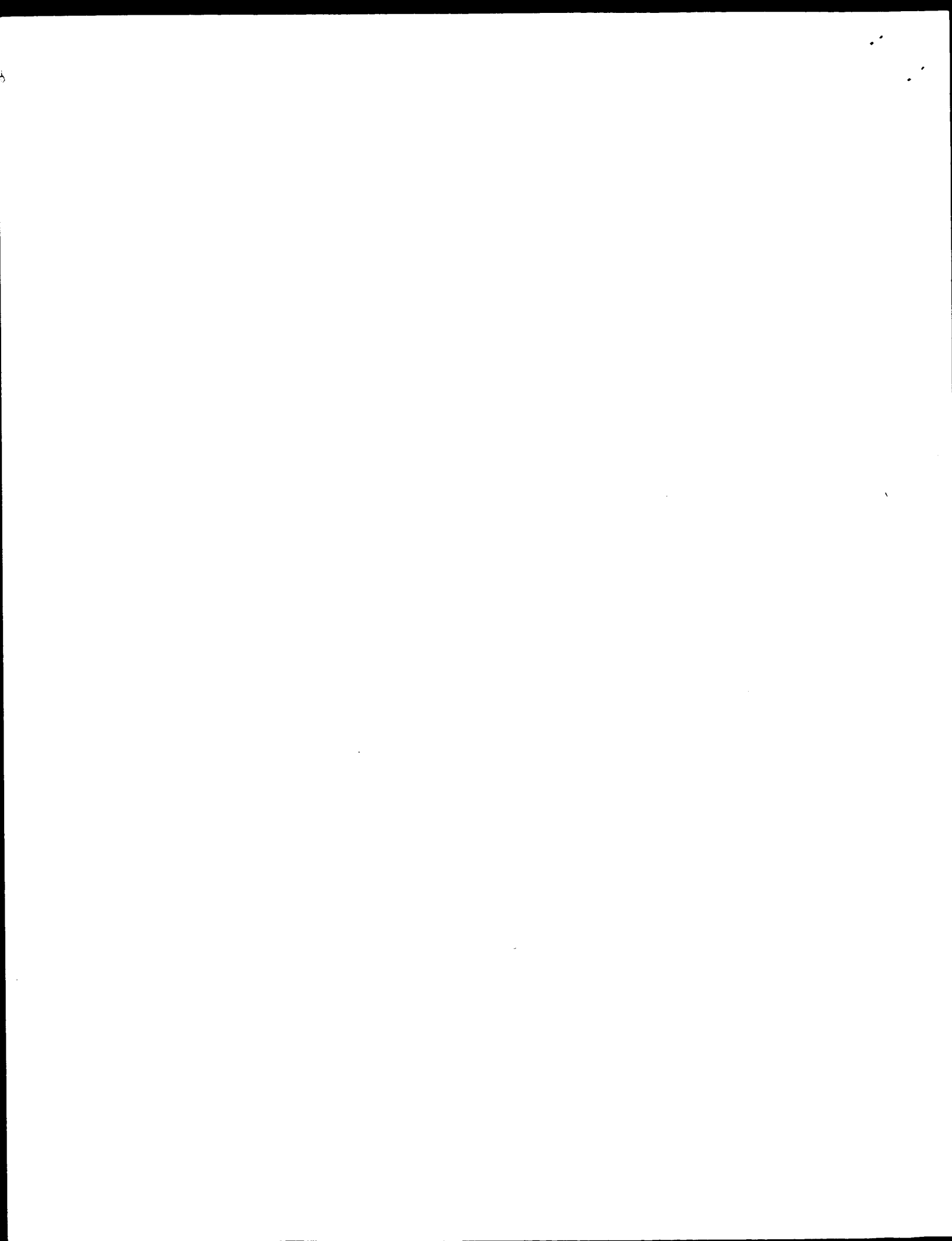
In another specific embodiment, the present invention is  
25 directed to the use of an agonist or antagonist of the IL-17 receptor like polypeptide, and/or an IL-17 receptor like polypeptide itself in combination (pretreatment, post-treatment, or concurrent treatment) with any of one or more slow-acting antirheumatic drugs (SAARDs) or disease modifying  
30 antirheumatic drugs (DMARDS), prodrug esters, or pharmaceutically acceptable salts thereof for the treatment of the diseases and disorders recited herein, including acute and chronic inflammation such as rheumatic diseases, graft versus host disease and multiple sclerosis. SAARDs or DMARDS,  
35 prodrug esters and pharmaceutically acceptable salts thereof

## FIGURE 2

Overlap of IL-17 Human Receptor-Like (hIL-17RL) Polypeptide  
and Known Human IL-17 Receptor Family Member

(SEQ ID NO: 3)

hIL-17RL	MMSGDWHDPGSQQCCLHLDSDMGLCVGRANWLHSASCSLVFPQGVGPASRN	50
Human IL-17R	..MGAARSPPS...AVPGPLLGLLLLLLGLVAPGGASLRLLDHRALVCSQ	45
hIL-17RL	SGLYNITFKYDNC..TTYLNVPVGKHVIADAQNITISQYACHDQ.....VA	93
Human IL-17R	PGLNCTVKNSTCLDDSWIHP.RNLTPSSPKDLQIQLFHAHTQQGDLFPV	93
hIL-17RL	VTILWS.PGALGIEFLKGFRVILEELKSEGRQCQQLILKDPKQLNSSFKR	142
Human IL-17R	AHIEWTLQTDASILYLEGAELSVLQNTNERLCVR..FEFLSKLRHHRR	141
hIL-17RL	TGMESQPFLNMKFETDYFVKV..VPFPSIKNESNYHPFFERTRACDLLLQ	190
Human IL-17R	WRFTFSHFV.VDPDQEYEVTVHHLPKPIPDGDPNHQSKNFLVPDCEHARM	190
hIL-17RL	PDNLACK...PFWKPR.....NL.NISQHGSDMQVSFDHA	221
Human IL-17R	KVTTPCMSSGSLWDPNITVETLEAHQLRVSFTLWNETHYQILLTSFPHM	240
hIL-17RL	PHNFGFRFFYLHYLKHGEPFKRKTCKQEQTTEMTSCLLQNVSPGDYIIE	271
Human IL-17R	ENHSCFEHMH.HIPAPRPEEFHQRSNVTLTLRNLKGCCRHOVQIQPFSS	289
hIL-17RL	LVDDT...NTTRKVMHYALKPVHSPWAGPIRAVAITVPLVVISAFATLFT	318
Human IL-17R	CLNDCLRHSATVSCPEMPDTPPEIPIDYMWVYWFITGISILLVGSVILL	339
hIL-17RL	VMCRKKQQENIYSHLDEESSESSTYTAALPRERLRPRP....KVFLCYSS	364
Human IL-17R	IVCMTWRLAGPGS...EKYSDDTKYTDGLPAADLIPPPLKPRKVIYISA	386
hIL-17RL	KDGQNHMNVVQCFAFYFLQDFCGCEVALDLWEDFSLCREGQREWV.IQK..	411
Human IL-17R	.DHPLYVDVVLKFAQFLLTACGTEVALDLLEEQAISEAGVMTWVGRQKQE	435
hIL-17RL	.IHESQFIIVVCSKGMK...YFVDKKNYKHGGGRGSGKGELFLVAVSA	456
Human IL-17R	MVESNSKIIVLCSRGRTRAKWQALLGRGAPVRLRCDHGKPVGDLFTAAMNM	485
hIL-17RL	IAEKLRLQAKQSSSAALSKFIAVYF.DYSCEGDVPGILDSTKYRLMDNLP	505
Human IL-17R	ILPDFKR.....PACFGTYVVCYFSEVSCDGDVPDLFGAAPRYPLMDRFE	530
hIL-17RL	QLCSHLHSRDHGLQEPGQHTROG..SRNRYFRSKSGRSLYVAICNMHQFI	553
Human IL-17R	EV..YFRIQDLEMFQPMHRVGEISGDNYLRSPGGRLRAALDRFRDWQ	578
hIL-17RL	DEEPDWFEKQFV....PFHPPLR...YREPVLKFDGSLVLNDVMCKPG	596
Human IL-17R	VRCPDWFECEENLYSADDQDAPSLDEEVFEPLPP.GTGIVKRAPLVRE.	626
hIL-17RL	PESDFCLKVEAAVLGATGPADSQHE.....SQHGG	626
Human IL-17R	PGSQACLAIDPLVGEEGGAATAKLEPHLQPRGQAPQPLHTLVLAEEGA	676





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comprise: allocupreide sodium, auranofin, aurothiogluco-  
se, aurothioglycanide, azathioprine, brequinar sodium,  
bucillamine, calcium 3-aurothio-2-propanol-1-sulfonate,  
chlorambucil, chloroquine, clobuzarit, cuproxoline,  
5 cyclophosphamide, cyclosporin, dapsone, 15-deoxyspergualin,  
diacerein, glucosamine, gold salts (e.g., cycloquine gold  
salt, gold sodium thiomalate, gold sodium thiosulfate),  
hydroxychloroquine, hydroxychloroquine sulfate, hydroxyurea,  
kebuzone, levamisole, lobenzarit, melittin, 6-mercaptapurine,  
10 methotrexate, mizoribine, mycophenolate mofetil, myoral,  
nitrogen mustard, D-penicillamine, pyridinol imidazoles such  
as SKNF86002 and SB203580, rapamycin, thiols, thymopoietin and  
vincristine. Structurally related SAARDs or DMARDs having  
similar analgesic and anti-inflammatory properties are also  
15 intended to be encompassed by this group.

In another specific embodiment, the present invention is  
directed to the use of an agonist or antagonist of the IL-17  
receptor like polypeptide, and/or an IL-17 receptor like  
polypeptide itself in combination (pretreatment, post-  
20 treatment, or concurrent treatment) with any of one or more  
COX2 inhibitors, prodrug esters or pharmaceutically acceptable  
salts thereof for the treatment of the diseases and disorders  
recited herein, including acute and chronic inflammation.  
Examples of COX2 inhibitors, prodrug esters or  
25 pharmaceutically acceptable salts thereof include, for  
example, celecoxib. Structurally related COX2 inhibitors  
having similar analgesic and anti-inflammatory properties are  
also intended to be encompassed by this group.

In still another specific embodiment, the present  
30 invention is directed to the use of an agonist or antagonist  
of the IL-17 receptor like polypeptide, and/or an IL-17  
receptor like polypeptide itself in combination (pretreatment,  
post-treatment, or concurrent treatment) with any of one or  
more antimicrobials, prodrug esters or pharmaceutically  
35 acceptable salts thereof for the treatment of the diseases and

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disorders recited herein, including acute and chronic inflammation. Antimicrobials include, for example, the broad classes of penicillins, cephalosporins and other beta-lactams, aminoglycosides, azoles, quinolones, macrolides, rifamycins, 5 tetracyclines, sulfonamides, lincosamides and polymyxins. The penicillins include, but are not limited to penicillin G, penicillin V, methicillin, nafcillin, oxacillin, cloxacillin, dicloxacillin, floxacillin, ampicillin, ampicillin/sulbactam, amoxicillin, amoxicillin/clavulanate, hetacillin, cyclacillin, 10 bacampicillin, carbenicillin, carbenicillin indanyl, ticarcillin, ticarcillin/clavulanate, azlocillin, mezlocillin, peperacillin, and mecillinam. The cephalosporins and other beta-lactams include, but are not limited to cephalothin, cephapirin, cephalixin, cephradine, cefazolin, cefadroxil, 15 cefaclor, cefamandole, cefotetan, cefoxitin, ceruroxime, cefonicid, ceforadine, cefixime, cefotaxime, moxalactam, ceftizoxime, ceftriaxone, cephaloperazone, ceftazidime, imipenem and aztreonam. The aminoglycosides include, but are not limited to streptomycin, gentamicin, tobramycin, amikacin, 20 netilmicin, kanamycin and neomycin. The azoles include, but are not limited to fluconazole. The quinolones include, but are not limited to nalidixic acid, norfloxacin, enoxacin, ciprofloxacin, ofloxacin, sparfloxacin and temafloxacin. The macrolides include, but are not limited to erythromycin, 25 spiramycin and azithromycin. The rifamycins include, but are not limited to rifampin. The tetracyclines include, but are not limited to spicycline, chlortetracycline, clomocycline, demeclocycline, deoxycycline, guamecycline, lymecycline, meclocycline, methacycline, minocycline, oxytetracycline, 30 penimepicycline, pipacycline, rolitetracycline, sancycline, senociclin and tetracycline. The sulfonamides include, but are not limited to sulfanilamide, sulfamethoxazole, sulfacetamide, sulfadiazine, sulfisoxazole and co-trimoxazole (trimethoprim/sulfamethoxazole). The lincosamides include, 35 but are not limited to clindamycin and lincomycin. The

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polymyxins (polypeptides) include, but are not limited to polymyxin B and colistin.

### IL-17 Receptor Like Compositions and Administration

Therapeutic compositions are within the scope of the present invention. Such IL-17 receptor like pharmaceutical compositions may comprise a therapeutically effective amount of an IL-17 receptor like polypeptide or an IL-17 receptor like nucleic acid molecule in admixture with a pharmaceutically or physiologically acceptable formulation agent selected for suitability with the mode of administration. Pharmaceutical compositions may comprise a therapeutically effective amount of one or more IL-17 receptor like selective binding agents in admixture with a pharmaceutically or physiologically acceptable formulation agent selected for suitability with the mode of administration.

Acceptable formulation materials preferably are nontoxic to recipients at the dosages and concentrations employed.

The pharmaceutical composition may contain formulation materials for modifying, maintaining or preserving, for example, the pH, osmolarity, viscosity, clarity, color, isotonicity, odor, sterility, stability, rate of dissolution or release, adsorption or penetration of the composition. Suitable formulation materials include, but are not limited to, amino acids (such as glycine, glutamine, asparagine, arginine or lysine), antimicrobials, antioxidants (such as ascorbic acid, sodium sulfite or sodium hydrogen-sulfite), buffers (such as borate, bicarbonate, Tris-HCl, citrates, phosphates, other organic acids), bulking agents (such as mannitol or glycine), chelating agents (such as ethylenediamine tetraacetic acid (EDTA)), complexing agents

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(such as caffeine, polyvinylpyrrolidone, beta-cyclodextrin or hydroxypropyl-beta-cyclodextrin), fillers, monosaccharides, disaccharides, and other carbohydrates (such as glucose, mannose, or dextrans), proteins (such as serum albumin, gelatin or immunoglobulins), coloring, flavoring and diluting agents, emulsifying agents, hydrophilic polymers (such as polyvinylpyrrolidone), low molecular weight polypeptides, salt-forming counterions (such as sodium), preservatives (such as benzalkonium chloride, benzoic acid, salicylic acid, thimerosal, phenethyl alcohol, methylparaben, propylparaben, chlorhexidine, sorbic acid or hydrogen peroxide), solvents (such as glycerin, propylene glycol or polyethylene glycol), sugar alcohols (such as mannitol or sorbitol), suspending agents, surfactants or wetting agents (such as pluronics, PEG, sorbitan esters, polysorbates such as polysorbate 20, polysorbate 80, triton, tromethamine, lecithin, cholesterol, tyloxapal), stability enhancing agents (sucrose or sorbitol), tonicity enhancing agents (such as alkali metal halides (preferably sodium or potassium chloride), mannitol sorbitol), delivery vehicles, diluents, excipients and/or pharmaceutical adjuvants. See *Remington's Pharmaceutical Sciences*, 18<sup>th</sup> Ed., A.R. Gennaro, ed., Mack Publishing Company (1990).

The optimal pharmaceutical composition will be determined by one skilled in the art depending upon, for example, the intended route of administration, delivery format, and desired dosage. See for example, *Remington's Pharmaceutical Sciences*, *supra*. Such compositions may influence the physical state, stability, rate of *in vivo* release, and rate of *in vivo* clearance of the IL-17 receptor like molecule.

The primary vehicle or carrier in a pharmaceutical composition may be either aqueous or non-aqueous in nature. For example, a suitable vehicle or carrier may be water for injection, physiological saline solution, or artificial cerebrospinal fluid, possibly supplemented with other materials common in compositions for parenteral

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administration. Neutral buffered saline or saline mixed with serum albumin are further exemplary vehicles. Other exemplary pharmaceutical compositions comprise Tris buffer of about pH 7.0-8.5, or acetate buffer of about pH 4.0-5.5, which may further include sorbitol or a suitable substitute therefor. In one embodiment of the present invention, IL-17 receptor like polypeptide compositions may be prepared for storage by mixing the selected composition having the desired degree of purity with optional formulation agents (Remington's *Pharmaceutical Sciences, supra*) in the form of a lyophilized cake or an aqueous solution. Further, the IL-17 receptor like polypeptide product may be formulated as a lyophilizate using appropriate excipients such as sucrose.

The IL-17 receptor like pharmaceutical compositions can be selected for parenteral delivery. Alternatively, the compositions may be selected for inhalation or for delivery through the digestive tract, such as orally. The preparation of such pharmaceutically acceptable compositions is within the skill of the art.

The formulation components are present in concentrations that are acceptable to the site of administration. For example, buffers are used to maintain the composition at physiological pH or at slightly lower pH, typically within a pH range of from about 5 to about 8.

When parenteral administration is contemplated, the therapeutic compositions for use in this invention may be in the form of a pyrogen-free, parenterally acceptable aqueous solution comprising the desired IL-17 receptor like molecule in a pharmaceutically acceptable vehicle. A particularly suitable vehicle for parenteral injection is sterile distilled water in which a IL-17 receptor like molecule is formulated as a sterile, isotonic solution, properly preserved. Yet another preparation can involve the formulation of the desired molecule with an agent, such as injectable microspheres, bio-erodible particles, polymeric compounds (polylactic acid,

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polyglycolic acid), or beads, or liposomes, that provides for the controlled or sustained release of the product which may then be delivered as a depot injection. Hyaluronic acid may also be used, and this may have the effect of promoting sustained duration in the circulation. Other suitable means for the introduction of the desired molecule include implantable drug delivery devices.

Pharmaceutical compositions such as (1) slow-release formulations, (2) inhalant mists, or (3) orally active formulations are also envisioned. The IL-17 receptor like molecule pharmaceutical composition generally is formulated for parenteral administration. Such parenterally administered therapeutic compositions are typically in the form of a pyrogen-free, parenterally acceptable aqueous solution comprising the desired IL-17 receptor like molecule in a pharmaceutically acceptable vehicle. The IL-17 receptor like molecule pharmaceutical compositions also may include particulate preparations of polymeric compounds such as polylactic acid, polyglycolic acid, etc. or the introduction of the molecule into liposomes. Hyaluronic acid may also be used, and this may have the effect of promoting sustained duration in the circulation

In one embodiment, a pharmaceutical composition may be formulated for inhalation. For example, an IL-17 receptor like molecule may be formulated as a dry powder for inhalation. IL-17 receptor like polypeptide or IL-17 receptor like nucleic acid molecule inhalation solutions may also be formulated with a propellant for aerosol delivery. In yet another embodiment, solutions may be nebulized. Pulmonary administration is further described in PCT application no. PCT/US94/001875, which describes pulmonary delivery of chemically modified proteins.

It is also contemplated that certain formulations may be administered orally. In one embodiment of the present invention, IL-17 receptor like molecules which are

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administered in this fashion can be formulated with or without those carriers customarily used in the compounding of solid dosage forms such as tablets and capsules. For example, a capsule may be designed to release the active portion of the formulation at the point in the gastrointestinal tract when bioavailability is maximized and pre-systemic degradation is minimized. Additional agents can be included to facilitate absorption of the IL-17 receptor like molecule. Diluents, flavorings, low melting point waxes, vegetable oils, lubricants, suspending agents, tablet disintegrating agents, and binders may also be employed.

Another pharmaceutical composition may involve an effective quantity of IL-17 receptor like molecules in a mixture with non-toxic excipients which are suitable for the manufacture of tablets. By dissolving the tablets in sterile water, or other appropriate vehicle, solutions can be prepared in unit dose form. Suitable excipients include, but are not limited to, inert diluents, such as calcium carbonate, sodium carbonate or bicarbonate, lactose, or calcium phosphate; or binding agents, such as starch, gelatin, or acacia; or lubricating agents such as magnesium stearate, stearic acid, or talc.

Additional IL-17 receptor like pharmaceutical compositions will be evident to those skilled in the art, including formulations involving IL-17 receptor like polypeptides in sustained- or controlled-delivery formulations. Techniques for formulating a variety of other sustained- or controlled-delivery means, such as liposome carriers, bio-erodible microparticles or porous beads and depot injections, are also known to those skilled in the art. See for example, PCT/US93/00829 which describes controlled release of porous polymeric microparticles for the delivery of pharmaceutical compositions. Additional examples of sustained-release preparations include semipermeable polymer matrices in the form of shaped articles, e.g. films, or

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microcapsules. Sustained release matrices may include polyesters, hydrogels, polylactides (U.S. 3,773,919, EP 58,481), copolymers of L-glutamic acid and gamma ethyl-L-glutamate (Sidman et al., *Biopolymers*, 22:547-556 (1983)),  
5 poly (2-hydroxyethyl-methacrylate) (Langer et al., *J. Biomed. Mater. Res.*, 15:167-277 (1981) and Langer, *Chem. Tech.*, 12:98-105 (1982)), ethylene vinyl acetate (Langer et al., *supra*) or poly-D(-)-3-hydroxybutyric acid (EP 133,988). Sustained-release compositions also may include liposomes, which can be  
10 prepared by any of several methods known in the art. See e.g., Eppstein et al., *Proc. Natl. Acad. Sci. USA*, 82:3688-3692 (1985); EP 36,676; EP 88,046; EP 143,949.

The IL-17 receptor like pharmaceutical composition to be used for *in vivo* administration typically must be sterile.  
15 This may be accomplished by filtration through sterile filtration membranes. Where the composition is lyophilized, sterilization using these methods may be conducted either prior to, or following, lyophilization and reconstitution. The composition for parenteral administration may be stored in  
20 lyophilized form or in solution. In addition, parenteral compositions generally are placed into a container having a sterile access port, for example, an intravenous solution bag or vial having a stopper pierceable by a hypodermic injection needle.

25 Once the pharmaceutical composition has been formulated, it may be stored in sterile vials as a solution, suspension, gel, emulsion, solid, or a dehydrated or lyophilized powder. Such formulations may be stored either in a ready-to-use form or in a form (e.g., lyophilized) requiring reconstitution  
30 prior to administration.

In a specific embodiment, the present invention is directed to kits for producing a single-dose administration unit. The kits may each contain both a first container having a dried protein and a second container having an aqueous



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formulation. Also included within the scope of this invention are kits containing single and multi-chambered pre-filled syringes (e.g., liquid syringes and lyosyringes).

5 An effective amount of an IL-17 receptor like pharmaceutical composition to be employed therapeutically will depend, for example, upon the therapeutic context and objectives. One skilled in the art will appreciate that the appropriate dosage levels for treatment will thus vary depending, in part, upon the molecule delivered, the  
10 indication for which the IL-17 receptor like molecule is being used, the route of administration, and the size (body weight, body surface or organ size) and condition (the age and general health) of the patient. Accordingly, the clinician may titer the dosage and modify the route of administration to obtain  
15 the optimal therapeutic effect. A typical dosage may range from about 0.1  $\mu\text{g/kg}$  to up to about 100  $\text{mg/kg}$  or more, depending on the factors mentioned above. In other embodiments, the dosage may range from 0.1  $\mu\text{g/kg}$  up to about 100  $\text{mg/kg}$ ; or 1  $\mu\text{g/kg}$  up to about 100  $\text{mg/kg}$ ; or 5  $\mu\text{g/kg}$  up to  
20 about 100  $\text{mg/kg}$ .

The frequency of dosing will depend upon the pharmacokinetic parameters of the IL-17 receptor like molecule in the formulation used. Typically, a clinician will administer the composition until a dosage is reached that  
25 achieves the desired effect. The composition may therefore be administered as a single dose, or as two or more doses (which may or may not contain the same amount of the desired molecule) over time, or as a continuous infusion via implantation device or catheter. Further refinement of the  
30 appropriate dosage is routinely made by those of ordinary skill in the art and is within the ambit of tasks routinely performed by them. Appropriate dosages may be ascertained through use of appropriate dose-response data.

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The route of administration of the pharmaceutical composition is in accord with known methods, e.g. oral, inhalation, injection or infusion by intravenous, intraperitoneal, intracerebral (intra-parenchymal), intracerebroventricular, intramuscular, intra-ocular, intraarterial, intraportal, or intralesional routes, or by sustained release systems or implantation device. Where desired, the compositions may be administered continuously by infusion, by bolus injection or continuously by infusion, or by implantation device.

Alternatively or additionally, the composition may be administered locally via implantation into the affected area of a membrane, sponge, or other appropriate material on to which the desired molecule has been absorbed or encapsulated. Where an implantation device is used, the device may be implanted into any suitable tissue or organ, and delivery of the desired molecule may be via diffusion, timed release bolus, or continuous administration or via catheter using continuous infusion.

It will further be appreciated that the IL-17 receptor like polypeptides, including fragments, variants, and derivatives, may be employed alone, together, or in combination with other polypeptides and pharmaceutical compositions. For example, the IL-17 receptor like polypeptides may be used in combination with cytokines, growth factors, antibiotics, anti-inflammatories, and/or chemotherapeutic agents as is appropriate for the indication being treated.

In some cases, it may be desirable to use IL-17 receptor like pharmaceutical compositions in an ex vivo manner. In such instances, cells, tissues, or organs that have been removed from the patient are exposed to IL-17 receptor like pharmaceutical compositions after which the cells, tissues

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and/or organs are subsequently implanted back into the patient.

In other cases, an IL-17 receptor like polypeptide can be delivered by implanting certain cells that have been  
5 genetically engineered, using methods such as those described herein, to express and secrete the polypeptide. Such cells may be animal or human cells, and may be autologous, heterologous, or xenogeneic. Optionally, the cells may be immortalized. In order to decrease the chance of an  
10 immunological response, the cells may be encapsulated to avoid infiltration of surrounding tissues. The encapsulation materials are typically biocompatible, semi-permeable polymeric enclosures or membranes that allow the release of the protein product(s) but prevent the destruction of the  
15 cells by the patient's immune system or by other detrimental factors from the surrounding tissues.

Additional embodiments of the present invention relate to cells and methods (e.g., homologous recombination and/or other recombinant production methods) for both the in vitro  
20 production of therapeutic polypeptides and for the production and delivery of therapeutic polypeptides by gene therapy or cell therapy. Homologous and other recombination methods may be used to modify a cell that contains a normally transcriptionally silent IL-17 receptor like gene, or an under  
25 expressed gene, and thereby produce a cell which expresses therapeutically efficacious amounts of IL-17 receptor like polypeptides.

It is further envisioned that IL-17 receptor like polypeptide can be produced in vitro or in vivo by homologous  
30 recombination, or with recombinant production methods utilizing control elements introduced into cells already containing DNA encoding IL-17 receptor like polypeptides. For example, homologous recombination is a technique originally developed for targeting genes to induce or correct mutations

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in transcriptionally active genes (Kucherlapati, *Prog. in Nucl. Acid Res. & Mol. Biol.*, 36:301, 1989). The basic technique was developed as a method for introducing specific mutations into specific regions of the mammalian genome (Thomas et al., *Cell*, 44:419-428, 1986; Thomas and Capecchi, *Cell*, 51:503-512, 1987; Doetschman et al., *Proc. Natl. Acad. Sci.*, 85:8583-8587, 1988) or to correct specific mutations within defective genes (Doetschman et al., *Nature*, 330:576-578, 1987). Exemplary homologous recombination techniques are described in U.S. Patent No. 5,272,071 (EP 9193051, EP Publication No. 505500; PCT/US90/07642, International Publication No. WO 91/09955).

Through homologous recombination, the DNA sequence to be inserted into the genome can be directed to a specific region of the gene of interest by attaching it to targeting DNA. The targeting DNA is a nucleotide sequence that is complementary (homologous) to a region of the genomic DNA. Small pieces of targeting DNA that are complementary to a specific region of the genome are put in contact with the parental strand during the DNA replication process. It is a general property of DNA that has been inserted into a cell to hybridize, and therefore, recombine with other pieces of endogenous DNA through shared homologous regions. If this complementary strand is attached to an oligonucleotide that contains a mutation or a different sequence or an additional nucleotide, it too is incorporated into the newly synthesized strand as a result of the recombination. As a result of the proofreading function, it is possible for the new sequence of DNA to serve as the template. Thus, the transferred DNA is incorporated into the genome.

Attached to these pieces of targeting DNA are regions of DNA which may interact with or control the expression of a IL-17 receptor like polypeptide, e.g., flanking sequences. For example, a promoter/enhancer element, a suppresser, or an

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exogenous transcription modulatory element is inserted in the genome of the intended host cell in proximity and orientation sufficient to influence the transcription of DNA encoding the desired IL-17 receptor like polypeptide. The control element  
5 controls a portion of the DNA present in the host cell genome. Thus, the expression of the desired IL-17 receptor like polypeptide may be achieved not by transfection of DNA that encodes the IL-17 receptor like gene itself, but rather by the use of targeting DNA (containing regions of homology with the  
10 endogenous gene of interest) coupled with DNA regulatory segments that provide the endogenous gene sequence with recognizable signals for transcription of an IL-17 receptor like polypeptide.

In an exemplary method, the expression of a desired  
15 targeted gene in a cell (*i.e.*, a desired endogenous cellular gene) is altered via homologous recombination into the cellular genome at a preselected site, by the introduction of DNA which includes at least a regulatory sequence, an exon and a splice donor site. These components are introduced into the  
20 chromosomal (genomic) DNA in such a manner that this, in effect, results in the production of a new transcription unit (in which the regulatory sequence, the exon and the splice donor site present in the DNA construct are operatively linked to the endogenous gene). As a result of the introduction of  
25 these components into the chromosomal DNA, the expression of the desired endogenous gene is altered.

Altered gene expression, as described herein, encompasses activating (or causing to be expressed) a gene which is normally silent (unexpressed) in the cell as obtained, as well  
30 as increasing the expression of a gene which is not expressed at physiologically significant levels in the cell as obtained. The embodiments further encompass changing the pattern of regulation or induction such that it is different from the pattern of regulation or induction that occurs in the cell as

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obtained, and reducing (including eliminating) the expression of a gene which is expressed in the cell as obtained.

One method by which homologous recombination can be used to increase, or cause, IL-17 receptor like polypeptide production from a cell's endogenous IL-17 receptor like gene involves first using homologous recombination to place a recombination sequence from a site-specific recombination system (e.g., Cre/loxP, FLP/FRT) (Sauer, *Current Opinion In Biotechnology*, 5:521-527, 1994; Sauer, *Methods In Enzymology*, 225:890-900, 1993) upstream (that is, 5' to) of the cell's endogenous genomic IL-17 receptor like polypeptide coding region. A plasmid containing a recombination site homologous to the site that was placed just upstream of the genomic IL-17 receptor like polypeptide coding region is introduced into the modified cell line along with the appropriate recombinase enzyme. This recombinase causes the plasmid to integrate, via the plasmid's recombination site, into the recombination site located just upstream of the genomic IL-17 receptor like polypeptide coding region in the cell line (Baubonis and Sauer, *Nucleic Acids Res.*, 21:2025-2029, 1993; O'Gorman et al., *Science*, 251:1351-1355, 1991). Any flanking sequences known to increase transcription (e.g., enhancer/promoter, intron, translational enhancer), if properly positioned in this plasmid, would integrate in such a manner as to create a new or modified transcriptional unit resulting in *de novo* or increased IL-17 receptor like polypeptide production from the cell's endogenous IL-17 receptor like gene.

A further method to use the cell line in which the site specific recombination sequence had been placed just upstream of the cell's endogenous genomic IL-17 receptor like polypeptide coding region is to use homologous recombination to introduce a second recombination site elsewhere in the cell line's genome. The appropriate recombinase enzyme is then introduced into the two-recombination-site cell line, causing

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a recombination event (deletion, inversion, translocation) (Sauer, *Current Opinion In Biotechnology*, supra, 1994; Sauer, *Methods In Enzymology*, supra, 1993) that would create a new or modified transcriptional unit resulting in de novo or increased IL-17 receptor like polypeptide production from the cell's endogenous IL-17 receptor like gene.

An additional approach for increasing, or causing, the expression of IL-17 receptor like polypeptide from a cell's endogenous IL-17 receptor like gene involves increasing, or causing, the expression of a gene or genes (e.g., transcription factors) and/or decreasing the expression of a gene or genes (e.g., transcriptional repressors) in a manner which results in de novo or increased IL-17 receptor like polypeptide production from the cell's endogenous IL-17 receptor like gene. This method includes the introduction of a non-naturally occurring polypeptide (e.g., a polypeptide comprising a site specific DNA binding domain fused to a transcriptional factor domain) into the cell such that de novo or increased IL-17 receptor like polypeptide production from the cell's endogenous IL-17 receptor like gene results.

The present invention further relates to DNA constructs useful in the method of altering expression of a target gene. In certain embodiments, the exemplary DNA constructs comprise: (a) one or more targeting sequences; (b) a regulatory sequence; (c) an exon; and (d) an unpaired splice-donor site. The targeting sequence in the DNA construct directs the integration of elements (a)-(d) into a target gene in a cell such that the elements (b)-(d) are operatively linked to sequences of the endogenous target gene. In another embodiment, the DNA constructs comprise: (a) one or more targeting sequences, (b) a regulatory sequence, (c) an exon, (d) a splice-donor site, (e) an intron, and (f) a splice-acceptor site, wherein the targeting sequence directs the integration of elements (a)-(f) such that the elements of (b)-

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(f) are operatively linked to the endogenous gene. The targeting sequence is homologous to the preselected site in the cellular chromosomal DNA with which homologous recombination is to occur. In the construct, the exon is generally 3' of the regulatory sequence and the splice-donor site is 3' of the exon.

If the sequence of a particular gene is known, such as the nucleic acid sequence of IL-17 receptor like polypeptide presented herein, a piece of DNA that is complementary to a selected region of the gene can be synthesized or otherwise obtained, such as by appropriate restriction of the native DNA at specific recognition sites bounding the region of interest. This piece serves as a targeting sequence(s) upon insertion into the cell and will hybridize to its homologous region within the genome. If this hybridization occurs during DNA replication, this piece of DNA, and any additional sequence attached thereto, will act as an Okazaki fragment and will be incorporated into the newly synthesized daughter strand of DNA. The present invention, therefore, includes nucleotides encoding a IL-17 receptor like polypeptide, which nucleotides may be used as targeting sequences.

IL-17 receptor like polypeptide cell therapy, e.g., the implantation of cells producing IL-17 receptor like polypeptides, is also contemplated. This embodiment involves implanting cells capable of synthesizing and secreting a biologically active form of IL-17 receptor like polypeptide. Such IL-17 receptor like polypeptide-producing cells can be cells that are natural producers of IL-17 receptor like polypeptides or may be recombinant cells whose ability to produce IL-17 receptor like polypeptides has been augmented by transformation with a gene encoding the desired IL-17 receptor like polypeptide or with a gene augmenting the expression of IL-17 receptor like polypeptide. Such a modification may be accomplished by means of a vector suitable for delivering the



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gene as well as promoting its expression and secretion. In order to minimize a potential immunological reaction in patients being administered an IL-17 receptor like polypeptide, as may occur with the administration of a polypeptide of a foreign species, it is preferred that the natural cells producing IL-17 receptor like polypeptide be of human origin and produce human IL-17 receptor like polypeptide. Likewise, it is preferred that the recombinant cells producing IL-17 receptor like polypeptide be transformed with an expression vector containing a gene encoding a human IL-17 receptor like polypeptide.

Implanted cells may be encapsulated to avoid the infiltration of surrounding tissue. Human or non-human animal cells may be implanted in patients in biocompatible, semipermeable polymeric enclosures or membranes that allow the release of IL-17 receptor like polypeptide, but that prevent the destruction of the cells by the patient's immune system or by other detrimental factors from the surrounding tissue. Alternatively, the patient's own cells, transformed to produce IL-17 receptor like polypeptides ex vivo, may be implanted directly into the patient without such encapsulation.

Techniques for the encapsulation of living cells are known in the art, and the preparation of the encapsulated cells and their implantation in patients may be routinely accomplished. For example, Baetge et al. (WO95/05452; PCT/US94/09299) describe membrane capsules containing genetically engineered cells for the effective delivery of biologically active molecules. The capsules are biocompatible and are easily retrievable. The capsules encapsulate cells transfected with recombinant DNA molecules comprising DNA sequences coding for biologically active molecules operatively linked to promoters that are not subject to down regulation in vivo upon implantation into a mammalian host. The devices provide for the delivery of the molecules from living cells to

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specific sites within a recipient. In addition, see U.S. Patent Nos. 4,892,538, 5,011,472, and 5,106,627. A system for encapsulating living cells is described in PCT Application no. PCT/US91/00157 of Aebischer et al. See also, PCT Application  
5 no. PCT/US91/00155 of Aebischer et al., Winn et al., *Exper. Neurol.*, 113:322-329 (1991), Aebischer et al., *Exper. Neurol.*, 111:269-275 (1991); and Tresco et al., *ASAIO*, 38:17-23 (1992).

In vivo and in vitro gene therapy delivery of IL-17 receptor like polypeptides is also envisioned. One example of  
10 a gene therapy technique is to use the IL-17 receptor like gene (either genomic DNA, cDNA, and/or synthetic DNA) encoding a IL-17 receptor like polypeptide which may be operably linked to a constitutive or inducible promoter to form a "gene  
15 therapy DNA construct". The promoter may be homologous or heterologous to the endogenous IL-17 receptor like gene, provided that it is active in the cell or tissue type into which the construct will be inserted. Other components of the gene therapy DNA construct may optionally include, DNA  
20 molecules designed for site-specific integration (e.g., endogenous sequences useful for homologous recombination), tissue-specific promoter, enhancer(s) or silencer(s), DNA molecules capable of providing a selective advantage over the parent cell, DNA molecules useful as labels to identify transformed cells, negative selection systems, cell specific  
25 binding agents (as, for example, for cell targeting), cell-specific internalization factors, and transcription factors to enhance expression by a vector as well as factors to enable vector manufacture.

A gene therapy DNA construct can then be introduced into  
30 cells (either ex vivo or in vivo) using viral or non-viral vectors. One means for introducing the gene therapy DNA construct is by means of viral vectors as described herein. Certain vectors, such as retroviral vectors, will deliver the DNA construct to the chromosomal DNA of the cells, and the

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gene can integrate into the chromosomal DNA. Other vectors will function as episomes, and the gene therapy DNA construct will remain in the cytoplasm.

In yet other embodiments, regulatory elements can be included for the controlled expression of the IL-17 receptor like gene in the target cell. Such elements are turned on in response to an appropriate effector. In this way, a therapeutic polypeptide can be expressed when desired. One conventional control means involves the use of small molecule dimerizers or rapalogs (as described in WO9641865 (PCT/US96/099486); WO9731898. (PCT/US97/03137) and WO9731899 (PCT/US95/03157) used to dimerize chimeric proteins which contain a small molecule-binding domain and a domain capable of initiating biological process, such as a DNA-binding protein or transcriptional activation protein. The dimerization of the proteins can be used to initiate transcription of the transgene.

An alternative regulation technology uses a method of storing proteins expressed from the gene of interest inside the cell as an aggregate or cluster. The gene of interest is expressed as a fusion protein that includes a conditional aggregation domain which results in the retention of the aggregated protein in the endoplasmic reticulum. The stored proteins are stable and inactive inside the cell. The proteins can be released, however, by administering a drug (e.g., small molecule ligand) that removes the conditional aggregation domain and thereby specifically breaks apart the aggregates or clusters so that the proteins may be secreted from the cell. See, *Science* 287:816-817, and 826-830 (2000).

Other suitable control means or gene switches include, but are not limited to, the following systems. Mifepristone (RU486) is used as a progesterone antagonist. The binding of a modified progesterone receptor ligand-binding domain to the

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progesterone antagonist activates transcription by forming a dimer of two transcription factors which then pass into the nucleus to bind DNA. The ligand binding domain is modified to eliminate the ability of the receptor to bind to the natural  
5 ligand. The modified steroid hormone receptor system is further described in U.S. 5,364,791; WO9640911, and WO9710337.

Yet another control system uses ecdysone (a fruit fly steroid hormone) which binds to and activates an ecdysone receptor (cytoplasmic receptor). The receptor then  
10 translocates to the nucleus to bind a specific DNA response element (promoter from ecdysone-responsive gene). The ecdysone receptor includes a transactivation domain/DNA-binding domain/ligand-binding domain to initiate transcription. The ecdysone system is further described in  
15 U.S. 5,514,578; WO9738117; WO9637609; and WO9303162.

Another control means uses a positive tetracycline-controllable transactivator. This system involves a mutated tet repressor protein DNA-binding domain (mutated tet R-4 amino acid changes which resulted in a reverse tetracycline-  
20 regulated transactivator protein, i.e., it binds to a tet operator in the presence of tetracycline) linked to a polypeptide which activates transcription. Such systems are described in U.S. Patent Nos. 5,464,758; 5,650,298 and 5,654,168.

25 Additional expression control systems and nucleic acid constructs are described in U.S. Patent Nos. 5,741,679 and 5,834,186, to Innovir Laboratories Inc.

In vivo gene therapy may be accomplished by introducing the gene encoding an IL-17 receptor like polypeptide into  
30 cells via local injection of an IL-17 receptor like nucleic acid molecule or by other appropriate viral or non-viral delivery vectors. Hefti, *Neurobiology*, 25:1418-1435 (1994). For example, a nucleic acid molecule encoding an IL-17

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receptor like polypeptide may be contained in an adeno-associated virus (AAV) vector for delivery to the targeted cells (e.g., Johnson, International Publication No. WO95/34670; International Application No. PCT/US95/07178).

- 5 The recombinant AAV genome typically contains AAV inverted terminal repeats flanking a DNA sequence encoding an IL-17 receptor like polypeptide operably linked to functional promoter and polyadenylation sequences.

- Alternative suitable viral vectors include, but are not  
10 limited to, retrovirus, adenovirus, herpes simplex virus, lentivirus, hepatitis virus, parvovirus, papovavirus, poxvirus, alphavirus, coronavirus, rhabdovirus, paramyxovirus, and papilloma virus vectors. U.S. Patent No. 5,672,344 describes an *in vivo* viral-mediated gene transfer system  
15 involving a recombinant neurotrophic HSV-1 vector. U.S. Patent No. 5,399,346 provides examples of a process for providing a patient with a therapeutic protein by the delivery of human cells which have been treated *in vitro* to insert a DNA segment encoding a therapeutic protein. Additional  
20 methods and materials for the practice of gene therapy techniques are described in U.S. Patent No. 5,631,236 involving adenoviral vectors; U.S. Patent No. 5,672,510 involving retroviral vectors; and U.S. 5,635,399 involving retroviral vectors expressing cytokines.

- 25 Nonviral delivery methods include, but are not limited to, liposome-mediated transfer, naked DNA delivery (direct injection), receptor-mediated transfer (ligand-DNA complex), electroporation, calcium phosphate precipitation, and microparticle bombardment (e.g., gene gun). Gene therapy  
30 materials and methods may also include the use of inducible promoters, tissue-specific enhancer-promoters, DNA sequences designed for site-specific integration, DNA sequences capable of providing a selective advantage over the parent cell, labels to identify transformed cells, negative selection

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systems and expression control systems (safety measures), cell-specific binding agents (for cell targeting), cell-specific internalization factors, and transcription factors to enhance expression by a vector as well as methods of vector  
5 manufacture. Such additional methods and materials for the practice of gene therapy techniques are described in U.S. Patent No. 4,970,154 involving electroporation techniques; WO96/40958 involving nuclear ligands; U.S. Patent No. 5,679,559 describing a lipoprotein-containing system for gene  
10 delivery; U.S. Patent No. 5,676,954 involving liposome carriers; U.S. Patent No. 5,593,875 concerning methods for calcium phosphate transfection; and U.S. Patent No. 4,945,050 wherein biologically active particles are propelled at cells at a speed whereby the particles penetrate the surface of the  
15 cells and become incorporated into the interior of the cells.

It is also contemplated that IL-17 receptor like gene therapy or cell therapy can further include the delivery of one or more additional polypeptide(s) in the same or a different cell(s). Such cells may be separately introduced  
20 into the patient, or the cells may be contained in a single implantable device, such as the encapsulating membrane described above, or the cells may be separately modified by means of viral vectors.

A means to increase endogenous IL-17 receptor like polypeptide expression in a cell via gene therapy is to insert  
25 one or more enhancer elements into the IL-17 receptor like polypeptide promoter, where the enhancer element(s) can serve to increase transcriptional activity of the IL-17 receptor like gene. The enhancer element(s) used will be selected  
30 based on the tissue in which one desires to activate the gene(s); enhancer elements known to confer promoter activation in that tissue will be selected. For example, if a gene encoding a IL-17 receptor like polypeptide is to be "turned on" in T-cells, the *lck* promoter enhancer element may be used.

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appropriate cells (either *ex vivo* or *in vivo*) either directly or via a viral vector as described herein. Typically, the integration of the construct into the genomic DNA of the cells will be via homologous recombination, where the 5' and 3' DNA sequences in the promoter construct can serve to help integrate the modified promoter region via hybridization to the endogenous chromosomal DNA.

#### Additional Uses of IL-17 receptor like Nucleic Acids and Polypeptides

Nucleic acid molecules of the present invention (including those that do not themselves encode biologically active polypeptides) may be used to map the locations of the IL-17 receptor like gene and related genes on chromosomes. Mapping may be done by techniques known in the art, such as PCR amplification and *in situ* hybridization.

IL-17 receptor like nucleic acid molecules (including those that do not themselves encode biologically active polypeptides), may be useful as hybridization probes in diagnostic assays to test, either qualitatively or quantitatively, for the presence of an IL-17 receptor like DNA or corresponding RNA in mammalian tissue or bodily fluid samples.

The IL-17 receptor like polypeptides may be used (simultaneously or sequentially) in combination with one or more cytokines, growth factors, antibiotics, anti-inflammatories, and/or chemotherapeutic agents as is appropriate for the indication being treated.

Other methods may also be employed where it is desirable to inhibit the activity of one or more IL-17 receptor like polypeptides. Such inhibition may be effected by nucleic acid molecules which are complementary to and hybridize to expression control sequences (triple helix formation) or to IL-17 receptor like mRNA. For example, antisense DNA or RNA

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Here, the functional portion of the transcriptional element to be added may be inserted into a fragment of DNA containing the IL-17 receptor like polypeptide promoter (and optionally, inserted into a vector and/or 5' and/or 3' flanking sequence(s), etc.) using standard cloning techniques. This construct, known as a "homologous recombination construct", can then be introduced into the desired cells either *ex vivo* or *in vivo*.

Gene therapy also can be used to decrease IL-17 receptor like polypeptide expression by modifying the nucleotide sequence of the endogenous promoter(s). Such modification is typically accomplished via homologous recombination methods. For example, a DNA molecule containing all or a portion of the promoter of the IL-17 receptor like gene(s) selected for inactivation can be engineered to remove and/or replace pieces of the promoter that regulate transcription. For example the TATA box and/or the binding site of a transcriptional activator of the promoter may be deleted using standard molecular biology techniques; such deletion can inhibit promoter activity thereby repressing the transcription of the corresponding IL-17 receptor like gene. The deletion of the TATA box or the transcription activator binding site in the promoter may be accomplished by generating a DNA construct comprising all or the relevant portion of the IL-17 receptor like polypeptide promoter(s) (from the same or a related species as the IL-17 receptor like gene(s) to be regulated) in which one or more of the TATA box and/or transcriptional activator binding site nucleotides are mutated via substitution, deletion and/or insertion of one or more nucleotides. As a result, the TATA box and/or activator binding site has decreased activity or is rendered completely inactive. The construct will typically contain at least about 500 bases of DNA that correspond to the native (endogenous) 5' and 3' DNA sequences adjacent to the promoter segment that has been modified. The construct may be introduced into the



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molecules, which have a sequence that is complementary to at least a portion of the selected IL-17 receptor like gene(s) can be introduced into the cell. Antisense probes may be designed by available techniques using the sequence of IL-17 receptor like polypeptide disclosed herein. Typically, each such antisense molecule will be complementary to the start site (5' end) of each selected IL-17 receptor like gene. When the antisense molecule then hybridizes to the corresponding IL-17 receptor like mRNA, translation of this mRNA is prevented or reduced. Antisense inhibitors provide information relating to the decrease or absence of an IL-17 receptor like polypeptide in a cell or organism.

Alternatively, gene therapy may be employed to create a dominant-negative inhibitor of one or more IL-17 receptor like polypeptides. In this situation, the DNA encoding a mutant polypeptide of each selected IL-17 receptor like polypeptide can be prepared and introduced into the cells of a patient using either viral or non-viral methods as described herein. Each such mutant is typically designed to compete with endogenous polypeptide in its biological role.

In addition, an IL-17 receptor like polypeptide, whether biologically active or not, may be used as an immunogen, that is, the polypeptide contains at least one epitope to which antibodies may be raised. Selective binding agents that bind to an IL-17 receptor like polypeptide (as described herein) may be used for *in vivo* and *in vitro* diagnostic purposes, including, but not limited to, use in labeled form to detect the presence of IL-17 receptor like polypeptide in a body fluid or cell sample. The antibodies may also be used to prevent, treat, or diagnose a number of diseases and disorders, including those recited herein. The antibodies may bind to an IL-17 receptor like polypeptide so as to diminish or block at least one activity characteristic of an IL-17 receptor like polypeptide, or may bind to a polypeptide to increase at least one activity characteristic of an IL-17

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receptor like polypeptide (including by increasing the pharmacokinetics of the IL-17 receptor like polypeptide).

#### EXAMPLE 1

##### 5 Cloning of cDNA Encoding human IL-17 Receptor Like Polypeptide

Materials and methods for cDNA cloning and analysis are described in Sambrook et al., supra. which is incorporated herein by reference.

10 An EST fragment zhgb-a1133097, was identified in an internal database. The fragment encoded a partial peptide sequence that is related to the human and mouse IL-17 receptor. The EST sequence was used to generate two gene specific primers, 2418-63 (5' CGA GCC ATG CTG GCT GAT GTT C 3': SEQ ID NO: 8) and 2418-64 (5' CGT GGT CGA AGG ACA CCT GCA 15 TG 3'; SEQ ID NO: 6), which correspond to the 5' region of the fragment. These primers were used to extend the 5' region of the fragment.

To isolate the full length cDNA, PCR was used to screen a 20 panel of 77 human tissue libraries. The PCR was carried out with Ready-To-Go Beads (Amersham Pharmacia Biotech, cat no. 27-9553-01), 50 ng of cDNA library and 5 pmol of each of the following primers: 2714-51 (5' CCA GTG TTT CGC CTA CTT CCT CC 3'; SEQ ID NO: 4) and 2417-56 (5' GAT ATC CGG TAA AGG GTT GGG 25 GC 3'; SEQ ID NO: 5). The PCR reactions were carried out at 94°C for 30 seconds, 5 cycles of 94°C for 15 seconds, 72°C for 2 minutes, then 5 cycles of 94°C for 15 seconds, 70°C for 2 minutes, followed by 25 cycles at 94°C for 15 seconds, 68°C for 2.5 minutes. Positive cDNA libraries expressed a 1247 bp 30 band. The following ten cDNA libraries were scored as positive: fetal pancreas random library, fetal kidney oligo-dT library, fetal ovary oligo-dT library, fetal calveria oligo-dT library, fetal femur oligo-dT library, fetal gall bladder oligo-dT library, fetal gall bladder random 35 library, spinal column oligo-dT library, spinal column random

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library, and bone (limb) oligo-dT library.

To further characterize the 10 positive libraries, 5'RACE PCR was carried out with 25 ng of cDNA from each library, 200  $\mu$ M dNTP, 1x Advantage cDNA Polymerase Mix (Clontech, CA cat. No. 8417-1) and 10 pmol of a gene specific primer 2418-64 (5' CGT GGT CGA AGG ACA CCT GCA TG 3'; SEQ ID NO: 6) and 10 pmol of a pSPORT vector primer 1916-83 (5' GGC TGT ATG TTG TGT GGA ATT GTG AGC G 3'; SEQ ID NO: 7) in a final volume of 50  $\mu$ l. The PCR reactions were carried out as follows: 94°C for 2 minutes, 5 cycles of 94°C for 15 seconds and 72°C for 4 minutes, then 5 cycles of 94°C for 15 seconds and 70°C for 4 minutes followed by 25 cycles of 94°C for 15 seconds and 68°C for 4 minutes.

The resulting PCR products were re-amplified using nested primers. Briefly, 5  $\mu$ l of 1:50 dilution of each product was used along with 10 pmol of the gene specific primer 2418-63 (5' CGA GCC ATG CTG GCT GAT GTT C 3'; SEQ ID NO: 8) and vector nest primer 1916-82 (5' CAT GAT TA CGC CAA GCT CTA ATA CGA CTC 3'; SEQ ID NO: 9) in each reaction. The RACE reactions were carried out as described above for the primary RACE reaction. The final RACE products (8  $\mu$ l) was analyzed on 1% TAE agarose gel at 5V/cm. The well-defined single bands were purified using the QIAquick PCR purification kit (QIAGEN, Cat. No. 28104) and sequenced using standard methods.

For the second round of 5' RACE, another gene specific primer 2432-38 (5' GAA GCT ACT GTT GAG CTG CTT CG 3'; SEQ ID NO: 10) and the pCMV/SPORT vector primer 2182-36 (5' CCG ATC CAG CCT CCG GAC TCT AG 3'; SEQ ID NO: 11) were used. Plasmid DNAs of LTI human multi-tissue cDNA library were used as templates. The PCR reactions were carried out in 50  $\mu$ l volume containing 30 ng of cDNA for each library pool, 10 pmol of each primer, 200  $\mu$ M dNTP and a 1x Advantage cDNA Polymerase Mix (Clontech, cat. No. 8417-1). The PCR reactions were carried out as follows: 94°C for 1 minute, followed by 35

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cycles of 94°C for 15 seconds, 65°C for 2 minutes and 72°C for 1 minute, followed by a final extension of 72°C for 10 minutes.

The resulting products were again re-amplified using another pair of nested primers, a gene specific primer 2144-06 (5' GCGTCAGCAATCACATGCTTCCC 3'; SEQ ID NO: 12) and the vector primer 2144-06 (5' GCCTATTTAGGTGACACTATAGAAC 3'; SEQ ID NO: 13). The PCR reactions were carried out as described above. The final products were analyzed by agarose gel electrophoresis, subcloned into the PCR 2.1-TOPO vector (Invitrogen Cat. No. K4560-01) and the inserts were sequenced using standard methods.

The resulting cDNA is set out as SEQ ID NO: 1 and is 3083 nucleotides in length. This cDNA contains a 2214 nucleotide coding region and encodes a 738 amino acid polypeptide which is set out as SEQ ID NO: 2 and is denoted as IL-17 receptor like polypeptide.

#### EXAMPLE 2

##### Tissue Expression of IL-17 Receptor like Polynucleotide

Quantitative PCR was carried out on various human fetal tissues to analyze the expression pattern of IL-17 receptor like mRNA. Total RNA was isolated with the Total RNA Isolation kit (Amersham Pharmacia Biotech, cat. No. 15593-031). The reverse transcriptase reactions were carried out as follows: 2 µg of total RNA was mixed with 1 µl (50 ng/µl) of random primer and incubated at 70°C for 10 minutes then quick-chilled on ice. Subsequently, 4 µl of 5x First Stand Buffer (BRL), 2 µl of 0.1 M DTT (BRL) and 1 µl of dNTP mix was added to the reactions and mixed well. The reactions were incubated at 37°C for 2 minutes. Subsequently, 1 µl of Superscript II reverse transcriptase (BRL) was added and incubated at 37°C, for 1 hour. The reactions were terminated by placing the tubes on ice.

The subsequent PCR reactions were carried using Ready-To-

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Go PCR Bead plates (Amersham Pharmacia Biotech cat. No. 27-9553-01). The following components were added to each well: 1  $\mu$ l of the RT reaction mixture after normalization with G#PDH levels, 1  $\mu$ l (10 pmol/ $\mu$ l) of primer 2417-51 (5' CCA GTG TTT CGC CTA CTT CCT CC 3'; SEQ ID NO: 14), 1  $\mu$ l (10 pmol/ $\mu$ l) of primer 2418-65 (5' GGA GCT TTT CGG CAA TGG CTG AC 3'; SEQ ID NO: 15) and 22  $\mu$ l of water. The reactions were mixed well carried out as follows: 94°C for 1 minute, followed by 5 cycles of 94°C for 30 seconds, 72°C for 4 minutes, 5 cycles of 94°C for 30 seconds, 70°C for 4 minutes and 25 cycles of 94°C for 30 seconds and 68°C for 4 minutes.

Expression of IL-17 receptor like transcript was detected on agarose gels and the intensities of expression as listed below. "++++" indicates a strong signal while "+" denotes a weak signal and "-" indicates no expression.

SOURCE	SIGNAL
Testis	++++
Kidney	++++
Intestine	+++
Pancreas	+++
Spinal Cord	+++
Bone	+++
Thymus	+
Placenta	-

### EXAMPLE 3

#### Production of IL-17 Receptor like Polypeptides

##### A. Bacterial Expression of IL-17 Receptor like Polypeptides

PCR is used to amplify template DNA sequences encoding a IL-17 receptor like polypeptide using primers corresponding to

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the 5' and 3' ends of the sequence. The amplified DNA products may be modified to contain restriction enzyme sites to allow for insertion into expression vectors. PCR products are gel purified and inserted into expression vectors using standard recombinant DNA methodology. An exemplary vector, such as pAMG21 (ATCC No. 98113) containing the lux promoter and a gene encoding kanamycin resistance is digested with BamHI and NdeI for directional cloning of inserted DNA. The ligated mixture is transformed into an *E. coli* host strain by electroporation and transformants are selected for kanamycin resistance. Plasmid DNA from selected colonies is isolated and subjected to DNA sequencing to confirm the presence of the insert.

Transformed host cells are incubated in 2x YT medium containing 30 mg/ml kanamycin at 30°C prior to induction.

Gene expression is induced by the addition of N-(3-oxohexanoyl)-dl-homoserine lactone to a final concentration of 30 ng/ml followed by incubation at either 30°C or 37°C for six hours. The expression of IL-17 receptor like polypeptide is evaluated by centrifugation of the culture, resuspension and lysis of the bacterial pellets, and analysis of host cell proteins by SDS-polyacrylamide gel electrophoresis.

Inclusion bodies containing IL-17 receptor like polypeptide are purified as follows. Bacterial cells are pelleted by centrifugation and resuspended in water. The cell suspension is lysed by sonication and pelleted by centrifugation at 195,000 x g for 5 to 10 minutes. The supernatant is discarded, and the pellet is washed and transferred to a homogenizer. The pellet is homogenized in 5 ml of a Percoll solution (75% liquid Percoll/0.15 M NaCl) until uniformly suspended and then diluted and centrifuged at 21,600 x g for 30 minutes. Gradient fractions containing the inclusion bodies are recovered and pooled. The isolated inclusion bodies are analyzed by SDS-PAGE. A single band on an SDS polyacrylamide gel corresponding to *E. coli*-produced IL-17 receptor like polypeptide is excised from the gel, and

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the N-terminal amino acid sequence is determined essentially as described by Matsudaira et al., J. Biol. Chem., 262: 10-35 (1987).

#### 5 B. Mammalian Cell Production of IL-17 Receptor like Polypeptides

The IL-17 receptor like DNA was subcloned into a mammalian expression vector as described above using standard DNA technology. An exemplary expression vector, pCEP4  
10 (Invitrogen, Carlsbad, CA), which contains an Epstein-Barr virus origin of replication, may be used for the expression of IL-17 receptor like in 293-EBNA-1 cells. Amplified and gel purified PCR products are ligated into pCEP4 vector and lipofected into 293-EBNA cells. The transfected cells are  
15 selected in 100 mg/ml hygromycin and the resulting drug-resistant cultures are grown to confluence. The cells are then cultured in serum-free media for 72 hours. The conditioned media is removed and, IL-17 receptor like protein polypeptide expression is analyzed by SDS-PAGE. Human E3 $\alpha$   
20 polypeptide expression may be detected by silver staining. Alternatively, IL-17 receptor like polypeptide is produced as a fusion protein with an epitope tag, such as an IgG constant domain or a FLAG epitope, which may be detected by Western blot analysis using antibodies to the tag peptide.

25 Human E3 $\alpha$  polypeptides may be excised from an SDS-polyacrylamide gel, or IL-17 receptor like fusion proteins are purified by affinity chromatography to the epitope tag, and subjected to N-terminal amino acid sequence analysis as described herein.

#### 30 EXAMPLE 4

##### Production of Anti- IL-17 receptor like Polypeptide

##### Antibodies

Polyclonal or monoclonal antibodies to IL-17 receptor  
35 like polypeptides may be obtained by immunization of animals

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with purified protein or with IL-17 receptor like peptides produced by biological or chemical synthesis. Suitable procedures for generating antibodies include those described in Hudson and Bay, Practical Immunology, Second Edition", .  
5 Edition, Blackwell Scientific Publications.

In one procedure for the production of monoclonal antibodies, animals (typically mice or rabbits) are injected with a IL-17 receptor like polypeptide antigens (such as an recombinant full length or truncated forms of IL-17 receptor  
10 like polypeptide, analogs, variants or the like), and those with sufficient serum titer levels as determined by ELISA are selected for hybridoma production. Spleens of immunized animals are collected and prepared as single cell suspensions from which splenocytes are recovered. The splenocytes are  
15 fused to mouse myeloma cells (such as Sp2/0-Ag14 cells), allowed to incubate in DMEM with 200 U/ml penicillin, 200 mg/ml streptomycin sulfate, and 4 mM glutamine, then incubated in HAT selection medium (Hypoxanthine; Aminopterin; Thymidine). After selection, the tissue culture supernatants  
20 are taken from each

fusion well and tested for anti- IL-17 receptor like polypeptide antibody production by ELISA.

Alternative procedures for obtaining anti- IL-17 receptor  
25 like polypeptide antibodies may also be employed, such as the immunization of transgenic mice harboring human Ig loci for production of human antibodies, and the screening of synthetic antibody libraries, such as those generated by mutagenesis of an antibody variable domain.

30



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WHAT IS CLAIMED

1. An isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:

(a) the nucleotide sequence as set forth in SEQ ID NO:

5 1;

(b) a nucleotide sequence encoding the polypeptide as set forth in SEQ ID NO: 2;

(c) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of

10 (a) or (b), wherein the encoded polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

(d) a nucleotide sequence complementary to any of (a)-(c).

15 2. An isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:

(a) a nucleotide sequence encoding a polypeptide that is at least about 70, 75, 80, 85, 90, 95, 96, 97, 98, or 99 percent identical to the polypeptide as set forth in SEQ ID  
20 NO: 2, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(b) a nucleotide sequence encoding an allelic variant or splice variant of the nucleotide sequence as set forth in SEQ ID NO: 1, wherein the encoded polypeptide has an activity of  
25 the polypeptide as set forth in SEQ ID NO: 2;

(c) a nucleotide sequence of SEQ ID NO: 1; (a); or (b) encoding a polypeptide fragment of at least about 25 amino acid residues, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

30 (d) a nucleotide sequence of SEQ ID NO: 1, or (a)-(c) comprising a fragment of at least about 16 nucleotides;

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(e) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of any of (a)-(d), wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

5 (f) a nucleotide sequence complementary to any of (a)-(c).

3. An isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:

10 (a) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one conservative amino acid substitution, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

15 (b) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one amino acid insertion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

20 (c) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one amino acid deletion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

25 (d) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 which has a C- and/or N- terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

30 (e) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 2 with at least one modification selected from the group consisting of amino acid substitutions, amino acid insertions, amino acid deletions, C-terminal truncation, and N-terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(f) a nucleotide sequence of (a)-(e) comprising a fragment of at least about 16 nucleotides;

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(g) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of any of (a)-(f), wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

5 (h) a nucleotide sequence complementary to any of (a)-(e).

4. A vector comprising the nucleic acid molecule of Claims 1, 2, or 3.

10

5. A host cell comprising the vector of Claim 4.

6. The host cell of Claim 5 that is a eukaryotic cell.

15 ~~7. The host cell of Claim 5 that is a prokaryotic cell.~~

8. A process of producing an IL-17 receptor like polypeptide comprising culturing the host cell of Claim 5 under suitable conditions to express the polypeptide, and  
20 optionally isolating the polypeptide from the culture.

9. A polypeptide produced by the process of Claim 8.

10. The process of Claim 8, wherein the nucleic acid  
25 molecule comprises promoter DNA other than the promoter DNA for the native IL-17 receptor like polypeptide operatively linked to the DNA encoding the IL-17 receptor like polypeptide.

30 11. The isolated nucleic acid molecule according to Claim 2 wherein the percent identity is determined using a computer program selected from the group consisting of GAP, BLASTP, BLASTN, FASTA, BLASTA, BLASTX, BestFit, and the Smith-Waterman algorithm.

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12. A process for determining whether a compound inhibits IL-17 receptor like polypeptide activity or production comprising exposing a cell according to Claims 5, 6, or 7 to the compound, and measuring IL-17 receptor like polypeptide activity or production in said cell.

13. An isolated polypeptide comprising the amino acid sequence set forth in SEQ ID NO: 2.

10 14. An isolated polypeptide comprising the amino acid sequence selected from the group consisting of:

(a) an amino acid sequence for an ortholog of SEQ ID NO: 2, wherein the encoded polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

15 (b) an amino acid sequence that is at least about 70, 80, 85, 90, 95, 96, 97, 98, or 99 percent identical to the amino acid sequence of SEQ ID NO: 2, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

20 (c) a fragment of the amino acid sequence set forth in SEQ ID NO: 2 comprising at least about 25 amino acid residues, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

25 (d) an amino acid sequence for an allelic variant or splice variant of either the amino acid sequence as set forth in SEQ ID NO: 2, or at least one of (a)-(b) wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2.

30 15. An isolated polypeptide comprising the amino acid sequence selected from the group consisting of:

(a) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one conservative amino acid substitution, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2 as determined using the computer

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program of GAP, BLASTP, BLASTN, FASTA, BLASTA, BLASTX, BestFit, or the Smith-Waterman algorithm,

(b) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one amino acid insertion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(c) the amino acid sequence as set forth in SEQ ID NO: 2 with at least one amino acid deletion, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2;

(d) the amino acid sequence as set forth in SEQ ID NO: 2 which has a C- and/or N-terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2; and

(e) the amino acid sequence as set forth in SEQ ID NO: 2, with at least one modification selected from the group consisting of amino acid substitutions, amino acid insertions, amino acid deletions, C-terminal truncation, and N-terminal truncation, wherein the polypeptide has an activity of the polypeptide as set forth in SEQ ID NO: 2.

16. An isolated polypeptide encoded by the nucleic acid molecule of Claims 1, 2, or 3.

17. The isolated polypeptide according to Claim 14 wherein the percent identity is determined using a computer program selected from the group consisting of GAP, BLASTP, BLASTN, FASTA, BLASTA, BLASTX, BestFit, and the Smith-Waterman algorithm.

18. A polypeptide according to claim 15 or 16 wherein the amino acid at position 45 of SEQ ID NO: 2 is glycine, proline, or alanine.

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19. A polypeptide according to claim 15 or 16 wherein the amino acid at position 227 of SEQ ID NO: 2 is phenylalanine, leucine, valine, isoleucine, alanine, or tyrosine.

5

20. A polypeptide according to claim 15 or 16 wherein the amino acid at position 363 of SEQ ID NO: 2 is serine, threonine, alanine or cysteine.

10 21. A polypeptide according to claim 15 or 16 wherein the amino acid at position 374 of SEQ ID NO: 2 is valine, isoleucine, methionine, leucine, phenylalanine, alanine or norleucine.

15 22. A polypeptide according to claim 15 or 16 wherein the amino acid at position 385 of SEQ ID NO: 2 is cysteine, serine, alanine.

20 23. A polypeptide according to claim 15 or 16 wherein the amino acid at position 515 of SEQ ID NO: 2 is aspartic acid or glutamic acid.

24. A polypeptide according to claim 15 or 16 wherein the amino acid at position 602 of SEQ ID NO: 2 is cysteine,  
25 alanine or serine.

25. An antibody produced by immunizing an animal with a peptide comprising an amino acid sequence of SEQ ID NO: 2.

30 26. An antibody or fragment thereof that specifically binds the polypeptide of Claims 13, 14, or 15.

27. The antibody of Claim 26 that is a monoclonal antibody.

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28. A hybridoma that produces a monoclonal antibody that binds to a peptide comprising an amino acid sequence of SEQ ID NO: 2.

5        29. A method of detecting or quantitating the amount of IL-17 receptor like polypeptide in a sample comprising contacting a sample suspected of containing IL-17 receptor like polypeptide with the anti-IL-17 receptor like antibody or fragment of Claims 25, 26, or 27 and detecting binding of said  
10 antibody or fragment.

30. A selective binding agent or fragment thereof that specifically binds at least one polypeptide wherein said polypeptide comprises the amino acid sequence selected from  
15 the group consisting of:

a) the amino acid sequence as set forth in SEQ ID NO: 2; and

b) a fragment of the amino acid sequence set forth in at least one of SEQ ID NO: 2; and

c) a naturally occurring variant thereof.

20

31. The selective binding agent of Claim 30 that is an antibody or fragment thereof.

32. The selective binding agent of Claim 30 that is a  
25 humanized antibody.

33. The selective binding agent of Claim 30 that is a human antibody or fragment thereof.

30        34. The selective binding agent of Claim 30 that is a polyclonal antibody or fragment thereof.

35. The selective binding agent Claim 30 that is a monoclonal antibody or fragment thereof.

35

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36. The selective binding agent of Claim 30 that is a chimeric antibody or fragment thereof.

37. The selective binding agent of Claim 30 that is a CDR-grafted antibody or fragment thereof.

38. The selective binding agent of Claim 30 that is an antiidiotypic antibody or fragment thereof.

39. The selective binding agent of Claim 30 which is a variable region fragment.

40. The variable region fragment of Claim 39 which is a Fab or a Fab' fragment.

41. A selective binding agent or fragment thereof comprising at least one complementarity determining region with specificity for a polypeptide having the amino acid sequence of SEQ ID NO: 2.

42. The selective binding agent of Claim 30 which is bound to a detectable label.

43. The selective binding agent of Claim 30 which antagonizes IL-17 receptor like polypeptide biological activity.

44. A method for treating, preventing, or ameliorating a disease, condition, or disorder associated with altered levels of IL-17 receptor like polypeptide comprising administering to a patient an effective amount of a selective binding agent according to Claim 30.

45. A selective binding agent produced by immunizing an animal with a polypeptide comprising an amino acid sequence



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selected from the group consisting of SEQ ID NO: 2.

46. A hybridoma that produces a selective binding agent capable of binding a polypeptide encoded by the nucleic acid of Claims 1, 2, or 3.

47. A composition comprising the polypeptide of Claims 13, 14, or 15 and a pharmaceutically acceptable formulation agent.

48. The composition of Claim 47 wherein the pharmaceutically acceptable formulation agent is a carrier, adjuvant, solubilizer, stabilizer, or anti-oxidant or combination thereof.

49. The composition of Claim 47 wherein the polypeptide comprises the mature amino acid sequence as set forth in SEQ ID NO: 2.

50. A polypeptide comprising a derivative of the polypeptide of Claims 13, 14, or 15.

51. The polypeptide of Claim 50 which is covalently modified with a water-soluble polymer.

52. The polypeptide of Claim 51 wherein the water-soluble polymer is selected from the group consisting of polyethylene glycol, monomethoxy-polyethylene glycol, dextran, cellulose, poly-(N-vinyl pyrrolidone) polyethylene glycol, propylene glycol homopolymers, polypropylene oxide/ethylene oxide copolymers, polyoxyethylated polyols, and polyvinyl alcohol.

53. A composition comprising a nucleic acid molecule of Claims 1, 2, or 3 and a pharmaceutically acceptable formulation agent.

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54. A composition of Claim 53 wherein said nucleic acid molecule is contained in a viral vector.

5

55. A viral vector comprising a nucleic acid molecule of Claims 1, 2, or 3.

56. A fusion polypeptide comprising the polypeptide of Claims 13, 14, or 15 fused to a heterologous amino acid sequence.

57. The fusion polypeptide of Claim 56 wherein the heterologous amino acid sequence is an IgG constant domain or fragment thereof.

58. A method for treating, preventing or ameliorating a medical condition comprising administering to a patient the polypeptide of Claims 13, 14, or 15 or the polypeptide encoded by the nucleic acid of Claims 1, 2, or 3 to said mammal.

59. A method of diagnosing a pathological condition or a susceptibility to a pathological condition in a subject caused by or resulting from abnormal levels of IL-17 receptor like polypeptide comprising:

(a) determining the presence or amount of expression of the polypeptide of Claims 13, 14, or 15 or the polypeptide encoded by the nucleic acid molecule of Claims 1, 2, or 3 in a sample; and

(b) diagnosing a pathological condition or a susceptibility to a pathological condition based on the presence or amount of expression of the polypeptide.

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60. A device, comprising:

(a) a membrane suitable for implantation; and

(b) cells encapsulated within said membrane, wherein said cells secrete a protein of Claims 13, 14, or 15, and wherein  
5 said membrane is permeable to said protein and impermeable to materials detrimental to said cells.

61. A device, comprising:

(a) a membrane suitable for implantation; and

10 (b) the IL-17 receptor like polypeptide encapsulated within said membrane, wherein said membrane is permeable to the polypeptide.

62. A method of identifying a compound which binds to a  
15 polypeptide comprising:

(a) contacting the polypeptide of Claims 13, 14, or 15 with a compound; and

(b) determining the extent of binding of the polypeptide to the compound.

20

63. A method of modulating levels of a polypeptide in an animal comprising administering to the animal the nucleic acid molecule of Claims 1, 2, or 3.

25 64. A transgenic non-human mammal comprising the nucleic acid molecule of Claims 1, 2, or 3.

65. A transgenic non-human comprising a disruption of the nucleic acid molecule of claim 1, 2 or 3 wherein the  
30 expression of IL-17 receptor polypeptide is decreased.

66. A method of identifying antagonists of IL-17 receptor like polypeptide biological activity comprising:

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(a) contacting a compound with an IL-17 receptor like polypeptide;

(b) detecting the biological activity of an IL-17 receptor like polypeptide in the presence of said compound;

5 and

(c) comparing the level of IL-17 receptor like polypeptide biological activity in the presence and absence of said compound.

10 67. Then method of claim 66 wherein the compound is a small molecule, peptide, protein, carbohydrate, or antibody.

68. A method of modulating levels of a polypeptide in an animal comprising administering to the animal the nucleic acid  
15 molecule of claims 1, 2, or 3.

69. An antagonist of IL-17 receptor like polypeptide activity selected from the group consisting of IL-17 receptor like selective binding agents, small molecules, antisense  
20 oligonucleotides, and peptides or derivatives thereof having specificity for IL-17 receptor like polypeptide.

70. A method of reducing cellular production of IL-17 receptor like polypeptide, comprising transforming or  
25 transfecting cells with a nucleic acid encoding an antagonist according to claim 69.

71. A method according to claim 70, wherein the antagonist is an antisense reagent, said reagent comprising an  
30 oligonucleotide comprising a single stranded nucleic acid sequence capable of binding to IL-17 receptor like mRNA.

FIGURE 1

(Map of Human IL-17 Receptor Like cDNA and Amino Acid Sequences)

(SEQ ID NO:1 and SEQ ID NO:2)

ATCATGATCCCTATGGCCTGGCAGTTAGGAATCTCCTGGGCTGGCGGGAGTTTGACAGCT 60

61 GTGGGGGGCCATGAAGCGGGCAGAAAGAGAGGTGGATGATGTCCGGGGACTGGCATGACC 120  
\* S G Q K E R W M M S G D W H D P -

121 CTGGGGTCACAGCAGTGCTGCTTGCTTGCATTTGGACTCCATGGGGCTTTGTGTTGGAAGAGCAA 180  
G S Q Q C C L H L D S M G L C V G R A N -

181 ATTGGCTTCACTCTGCATCATGTTCTCTTGTGTTTTCCACAGGGAGTGGGGCCAGCCAGCA 240  
W L H S A S C S L V F P Q G V G P A S R -

240 GAAACAGTGGGCTGTACAACATCACCTTCAAATATGACAATTGTACCACCTACTTGAATC 300  
N S G L Y N I T F K Y D N C T T Y L N P -

301 CAGTGGGGAAGCATGTGATTGCTGACGCCAGAATATCACCATCAGCCAGTATGCTTGCC 360  
V G K H V I A D A Q N I T I S Q Y A C H -

361 ATGACCAAGTGGCAGTCACCATTTCTTGGTCCCCAGGGGCCCTCGGCATCGAATTCCTGA 420  
D Q V A V T I L W S P G A L G I E F L K -

421 AAGGATTTTCGGGTAATACTGGAGGAGCTGAAGTCGAGGGAAGACAGTGCCAACAACCTGA 480  
G F R V I L E E L K S E G R Q C Q Q L I -

481 TTCTAAAGGATCCGAAGCAGCTCAACAGTAGCTTCAAAGAAGCTGGAATGGAATCTCAAC 540  
L K D P K Q L N S S F K R T G M E S Q P -

541 CTTTCCTGAATATGAAATTTGAAACGGATTATTTTCGTAAAGGTTGTCCCTTTTCCTTCCA 600  
F L N M K F E T D Y F V K V V P F P S I -

601 TTAAAAACGAAAGCAATTACCACCCTTTCTTCTTTAGAACCCGAGCCTGTGACCTGTTGT 660  
K N E S N Y H P F F F R T R A C D L L L -

661 TACAGCCGGACAATCTAGCTTGTAACCCTTCTGGAAGCCTCGGAACCTGAACATCAGCC 720  
Q P D N L A C K P F W K P R N L N I S Q -

721 AGCATGGCTCGGACATGCAGGTGTCCTTCGACCACGCACCGCACAACTTCGGCTTCCGTT 780  
H G S D M Q V S F D H A P H N F G F R F -

781 TCTTCTATCTTCACTACAAGCTCAAGCACGAAGGACCTTTCAAGCGAAAGACCTGTAAGC 840  
F Y L H Y K L K H E G P F K R K T C K Q -

841 AGGAGCAAACCTACAGAGATGACCAGCTGCCTCCTTCAAATGTTTCTCCAGGGGATTATA 900  
E Q T T E M T S C L L Q N V S P G D Y I -

901 TAATTGAGCTGGTGGATGACACTAACACAACAAGAAAGTGATGCATTATGCCTTAAAGC 960  
I E L V D D T N T T R K V M H Y A L K P -



3721 GGGGGGCTTGCACCCCTGCTGGAAAGAAAAGGTTTGTGTATATTTCTGATGCAAATGTCA 3780  
3781 TACTCACTGCTCTGTAAAGGCAGCTGGCAGCTTTTTGGGAAAAGAACGTGCTCGTCTGTT 3840  
3841 CTCTGGCATCAAGTTTCTTGCAGCTGCTCTGAGGGAGAGACAGTGAGCTGCAAGACTGCC 3900  
3901 TCCCCATAACAACAGGCAACTCAGAGAAGAGTCATTTTATGTTGTTTCTATGGAATCTGG 3960  
3961 AATGAGTGCAGAGCTCCTACCCACACATGACTGCCCCGCCATTTTCATCCTAGGCATTCTG 4020  
4021 TGAAGGAGATTGGTTAGTCCAAACTTGCTAACATACGAAAATTCAGTTGGAACATGATGA 4080  
4081 GAGATTTCTTATTGAGGCCAAGAGATGTTTCCTGTCCCAGAGGAACCATTAGGAGTCGCT 4140  
4141 TTTAGGGTATTCAGCTTTGTTCATGAAATAAGGCATCTCTGAGAAAGTGGCCCCAGGGAG 4200  
4201 AGAATGGAGGACTGGGAGGAGAAGCATTAACTGAGCTCCAAGGGTGTGTGGGCAGAGAGC 4260  
4261 TTGCTATGTGAACTCACTCCTTAAGAAAATGGAAGAGAAAAAGAGAGTGCTAGTTAAAAA 4320  
4321 ATCGGGATGTTTTAGTTTGGATTTAGGGTTTTGATACTTATGTTGAAATACTAATGTTTC 4380  
4381 TGATCAATAAAATCAAACCTCTTAATATACCGAGTAATGAAACCATAGTGTGATTGCCTCA 4440  
4441 GAATAAATTGAGAAGTCCAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 4483





hIL-17RL	LDQDGEARPALDGSAALQPLLHTVKA....GSPSDMPRDSGIYDSSVPSS	672
Human IL-17R	LVAAVEPGPLADGAAVRLALAGEGEACPLLGGPG.AGRNSVLF...LPVD	722
hIL-17RL	ELSLPLMEGLSTDQTETSSLTESVSSSSGLGEEPPALPSKLLSSGSCKA	722
Human IL-17R	PEDSPL..GSSTPMASPDLLPEDVR.....EHLEGLMLSLEQSLSCQA	764
hIL-17RL	DLGC.RSYTDELHAVAPL*	740
Human IL-17R	QGGCSRPMVLTDPHTPYEEERQSVQSDQGYISRSSPQPPEGLTEMEEE	814



-1-

## SEQUENCE LISTING

&lt;110&gt; Amgen Inc.

&lt;120&gt; IL-17 Receptor Like Molecules and Uses Thereof

&lt;130&gt; 01017/36916A

&lt;140&gt;

&lt;141&gt;

&lt;150&gt; 09/724,460

&lt;151&gt; 2000-11-28

&lt;150&gt; 60/189,816

&lt;151&gt; 2000-03-16

&lt;160&gt; 17

&lt;170&gt; PatentIn Ver. 2.0

&lt;210&gt; 1

&lt;211&gt; 3083

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

&lt;221&gt; CDS

&lt;222&gt; (22)..(2235)

&lt;400&gt; 1

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                        1                      5                      10

ttc ttt acg gtc aac gcc tgc ctc aac ggc tcg cag ctg gct gtg gcc      99
Phe Phe Thr Val Asn Ala Cys Leu Asn Gly Ser Gln Leu Ala Val Ala
                        15                      20                      25

gct ggc ggg tcc ggc cgc gcg tgg ggc gtc gac acc tgt ggc tgg agg     147
Ala Gly Gly Ser Gly Arg Ala Trp Gly Val Asp Thr Cys Gly Trp Arg
                        30                      35                      40

gga gtg ggg cca gcc agc aga aac agt ggg ctg tac aac atc acc ttc     195
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                        45                      50                      55

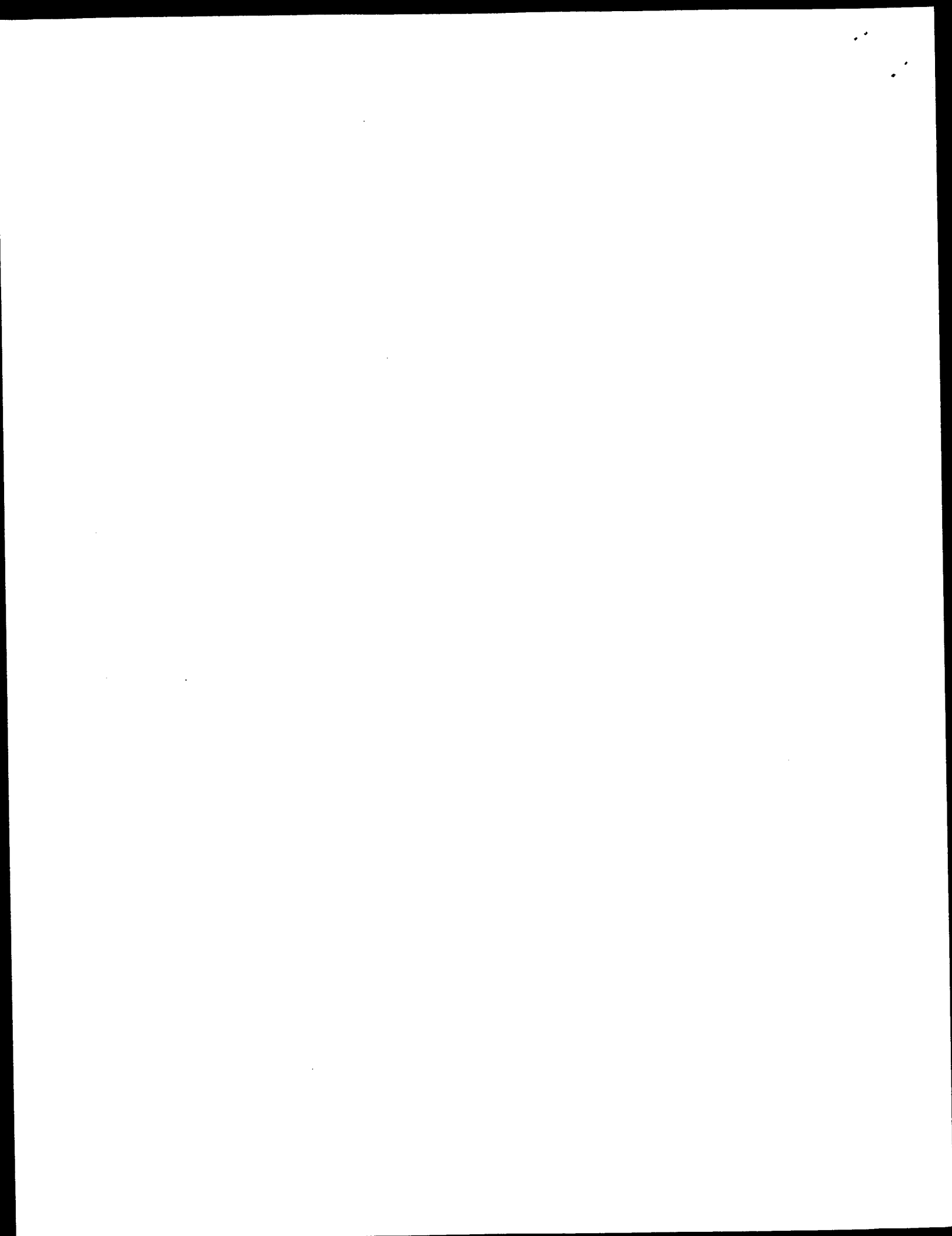
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Ile Ala Asp Ala Gln Asn Ile Thr Ile Ser Gln Tyr Ala Cys His Asp
                        75                      80                      85

caa gtg gca gtc acc att ctt tgg tcc cca ggg gcc ctc ggc atc gaa     339
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                        95                      100                      105

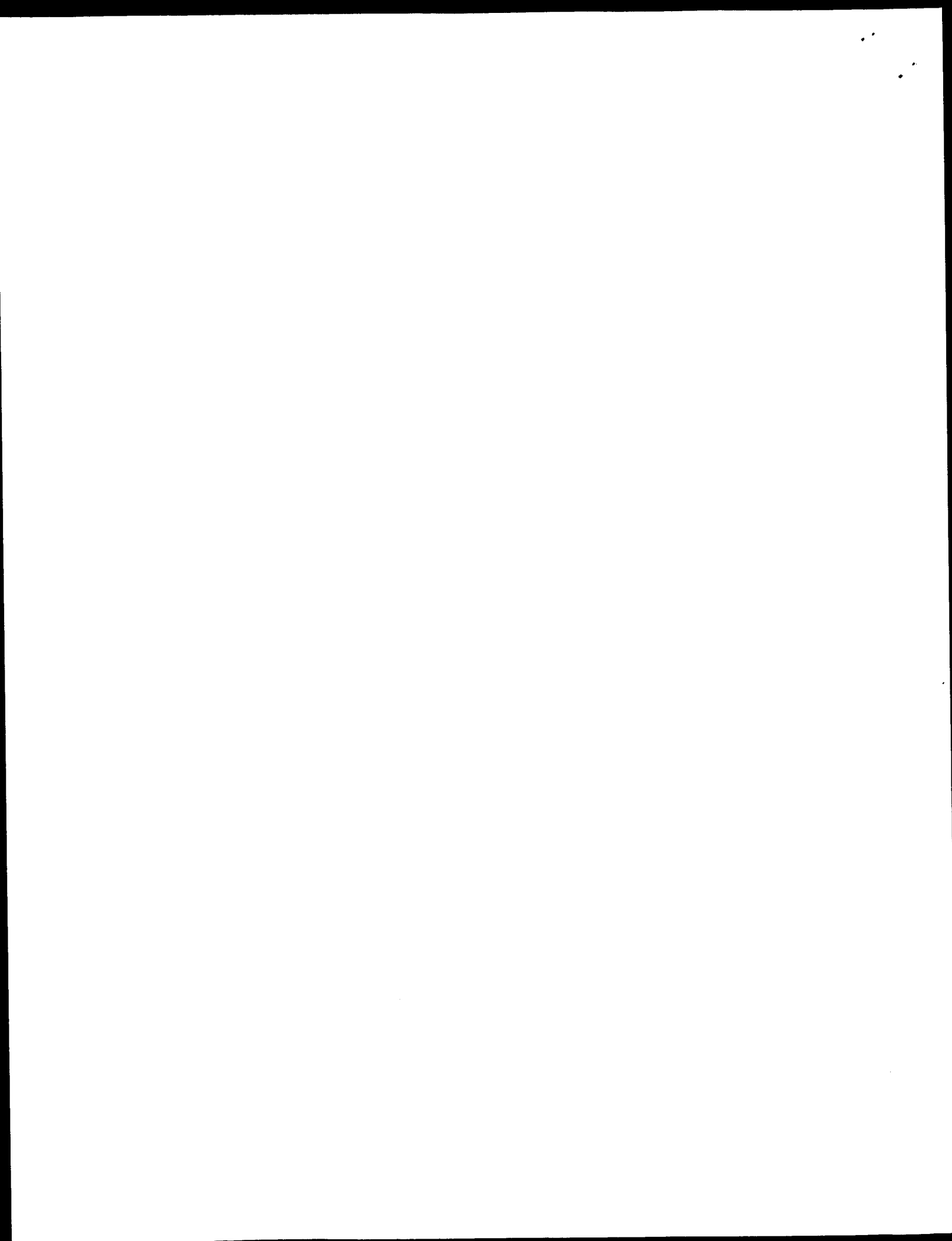
ttc ctg aaa gga ttt cgg gta ata ctg gag gag ctg aag tcg gag gga     387
Phe Leu Lys Gly Phe Arg Val Ile Leu Glu Glu Leu Lys Ser Glu Gly
                        110                      115                      120

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-2-

aga cag tgc caa caa ctg att cta aag gat ccg aag cag ctc aac agt	435
Arg Gln Cys Gln Gln Leu Ile Leu Lys Asp Pro Lys Gln Leu Asn Ser	
125 130 135	
agc ttc aaa aga act gga atg gaa tct caa cct ttc ctg aat atg aaa	483
Ser Phe Lys Arg Thr Gly Met Glu Ser Gln Pro Phe Leu Asn Met Lys	
140 145 150	
ttt gaa acg gat tat ttc gta aag gtt gtc cct ttt cct tcc att aaa	531
Phe Glu Thr Asp Tyr Phe Val Lys Val Val Pro Phe Pro Ser Ile Lys	
155 160 165 170	
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Asn Glu Ser Asn Tyr His Pro Phe Phe Phe Arg Thr Arg Ala Cys Asp	
175 180 185	
ctg ttg tta cag ccg gac aat cta gct tgt aaa ccc ttc tgg aag cct	627
Leu Leu Leu Gln Pro Asp Asn Leu Ala Cys Lys Pro Phe Trp Lys Pro	
190 195 200	
cgg aac ctg aac atc agc cag cat ggc tgc gac atg cag gtg tcc ttc	675
Arg Asn Leu Asn Ile Ser Gln His Gly Ser Asp Met Gln Val Ser Phe	
205 210 215	
gac cac gca ccg cac aac ttc ggc ttc cgt ttc ttc tat ctt cac tac	723
Asp His Ala Pro His Asn Phe Gly Phe Arg Phe Phe Tyr Leu His Tyr	
220 225 230	
aag ctc aag cac gaa gga cct ttc aag cga aag acc tgt aag cag gag	771
Lys Leu Lys His Glu Gly Pro Phe Lys Arg Lys Thr Cys Lys Gln Glu	
235 240 245 250	
caa act aca gag acg acc agc tgc ctc ctt caa aat gtt tct cca ggg	819
Gln Thr Thr Glu Thr Ser Cys Leu Leu Gln Asn Val Ser Pro Gly	
255 260 265	
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Asp Tyr Ile Ile Glu Leu Val Asp Asp Thr Asn Thr Thr Arg Lys Val	
270 275 280	
atg cat tat gcc tta aag cca gtg cac tcc ccg tgg gcc ggg ccc atc	915
Met His Tyr Ala Leu Lys Pro Val His Ser Pro Trp Ala Gly Pro Ile	
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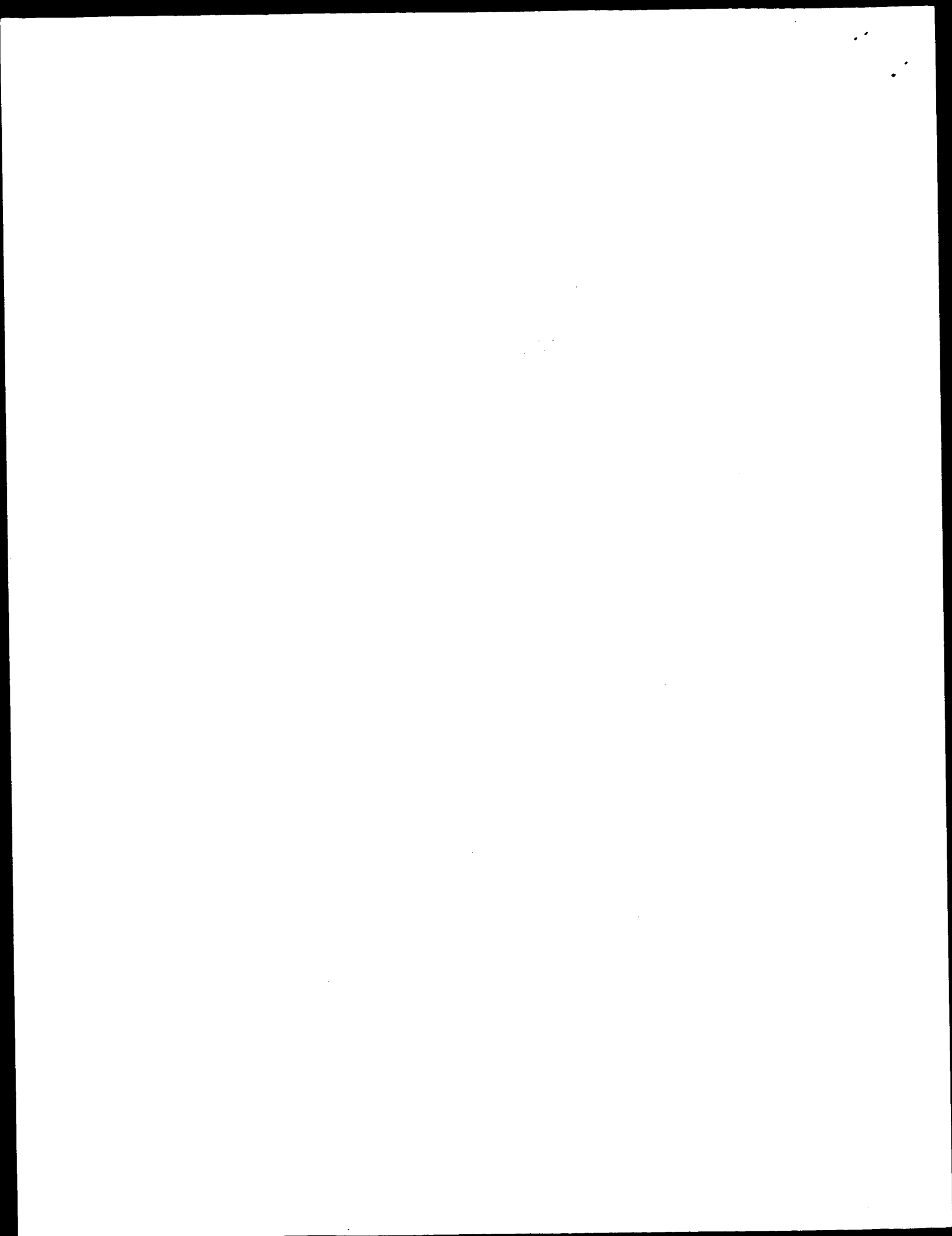
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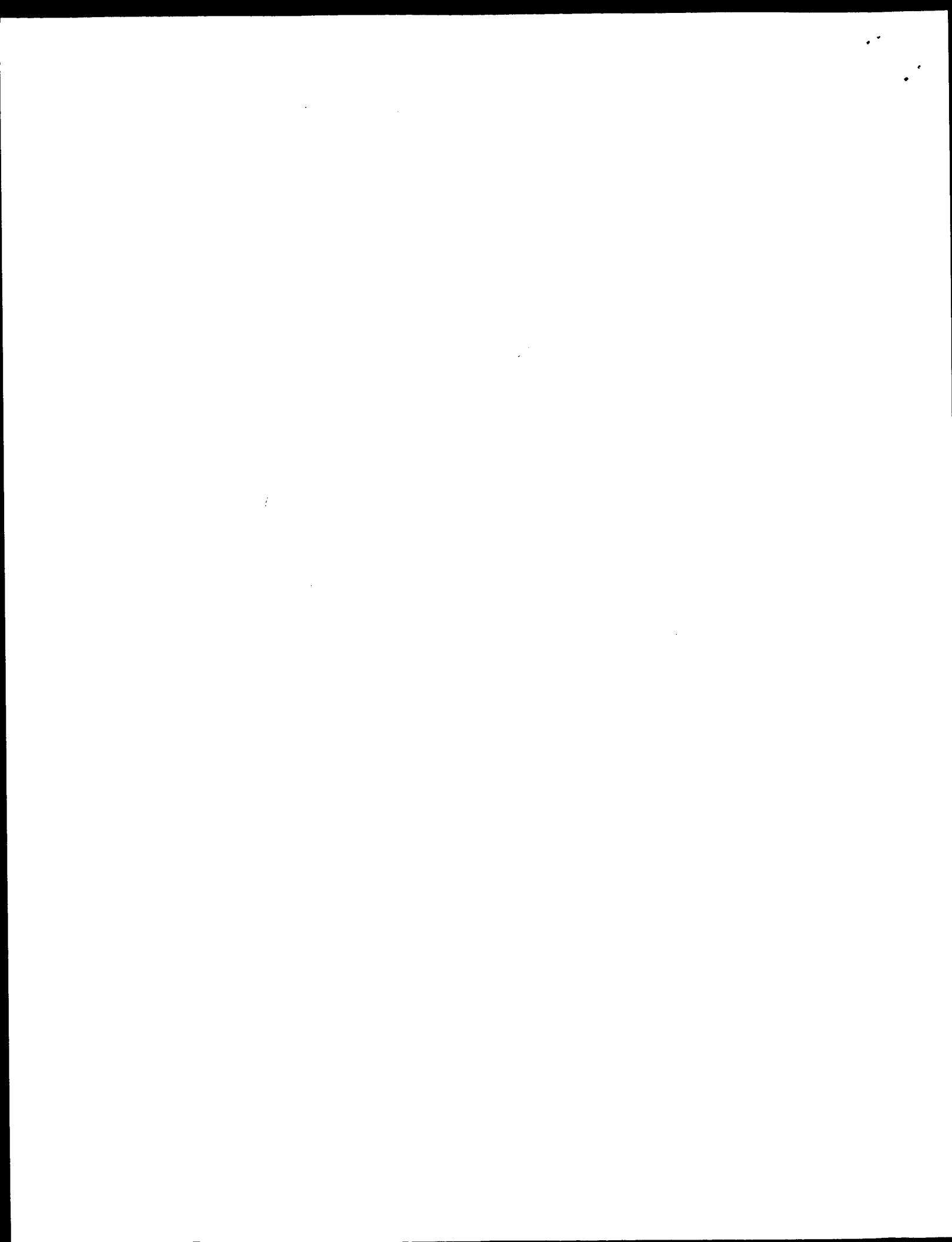
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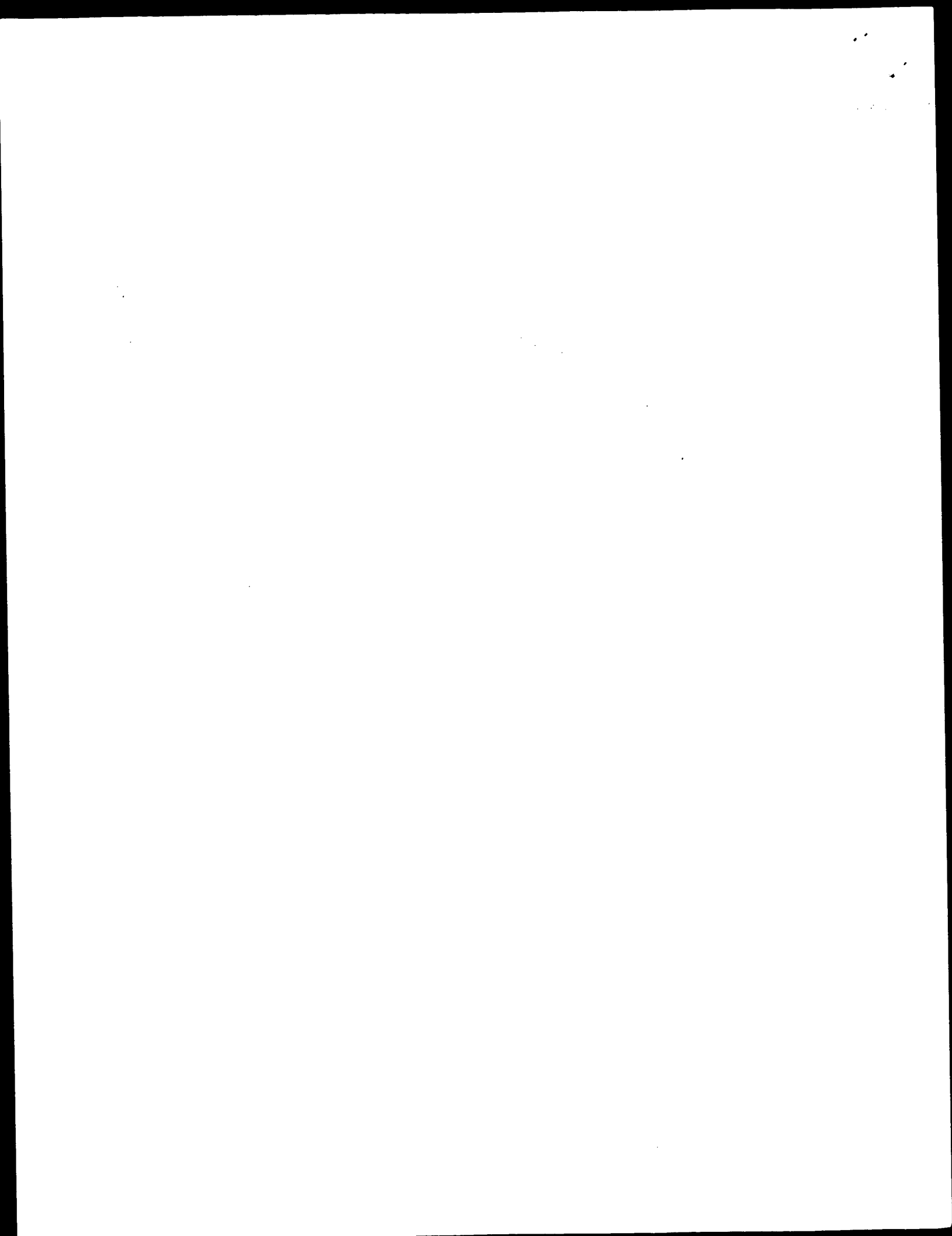
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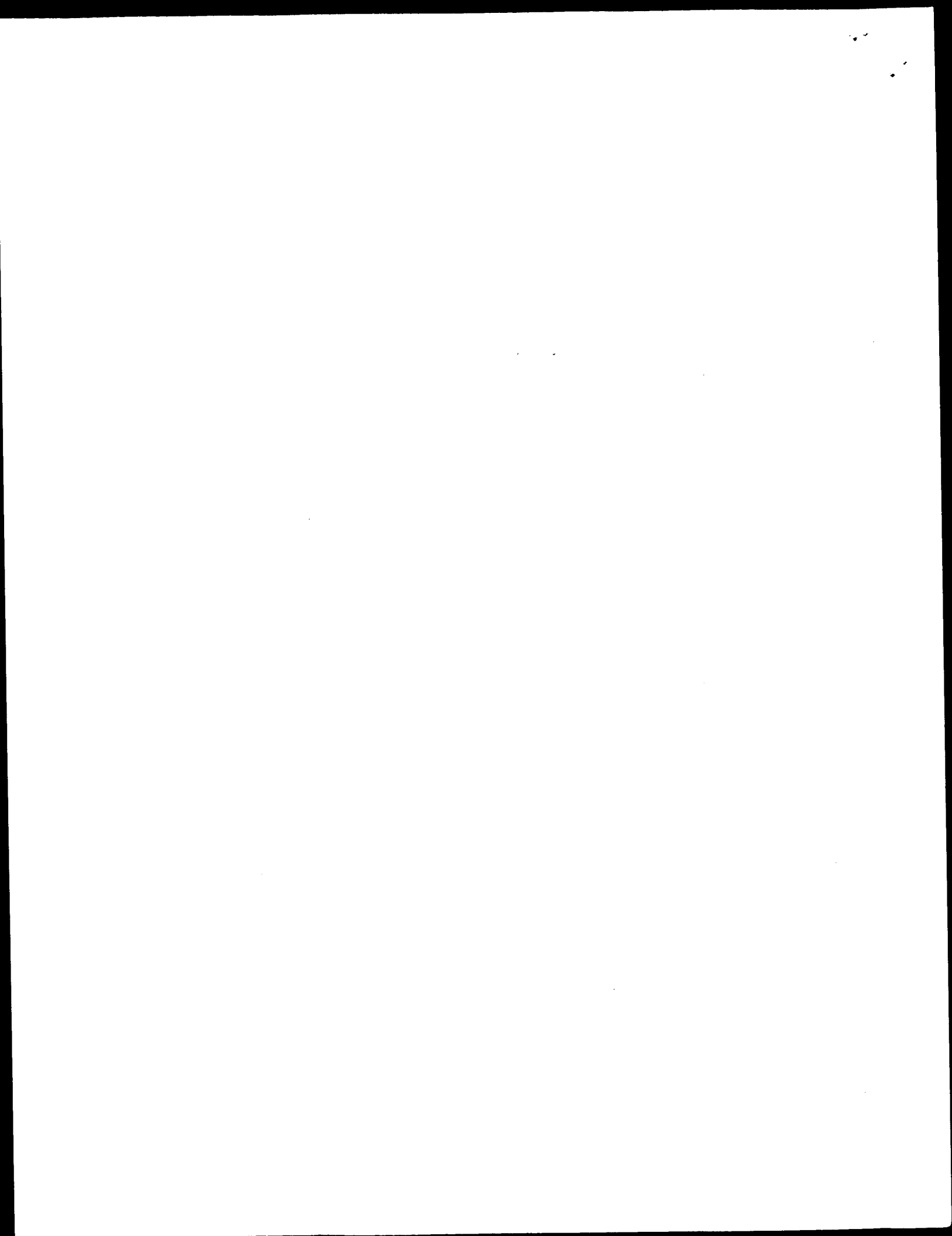
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(72) Inventor; and

(75) Inventor/Applicant (for US only): JING, Shugian  
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Wacker Drive, Chicago, IL 60606 (US).

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TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.

(84) Designated States (regional): ARIPO patent (GH, GM,  
KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian  
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CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

Published:

— with international search report

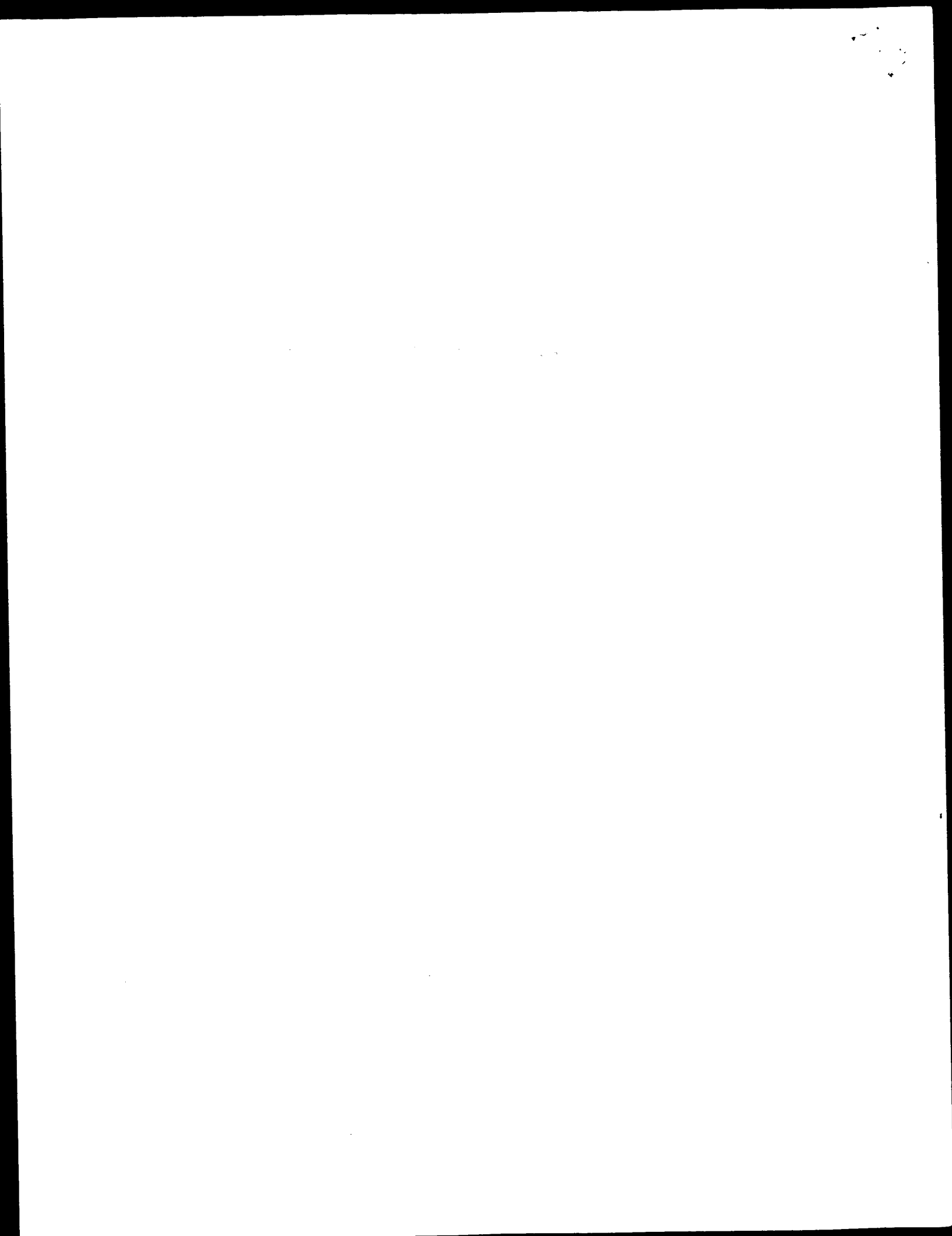
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For two-letter codes and other abbreviations, refer to the "Guid-  
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ning of each regular issue of the PCT Gazette.

(54) Title: IL-17 RECEPTOR LIKE MOLECULES AND USES THEREOF

(57) Abstract: Novel IL-17 receptor like polypeptides and nucleic acid molecules encoding the same. The invention also pro-  
vides vectors, host cells, agonists and antagonists (including selective binding agents), and method for producing IL-17 receptor like  
polypeptides. Also provided for are methods for the treatment, diagnosis, amelioration, or prevention of diseases with IL-17 receptor  
like polypeptides.

WO 01/68859 A3



# INTERNATIONAL SEARCH REPORT

Inte. onal Application No

PCT/US 01/08678

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	WO 96 29408 A (IMMUNEX CORP) 26 September 1996 (1996-09-26) the whole document especially seq ID no.9, no.10 and the examples claims ---	1-71
A	WO 99 14240 A (HUMAN GENOME SCIENCES INC ;RUBEN STEVEN M (US); SHI YANGGU (US)) 25 March 1999 (1999-03-25) the whole document ---	1-71
A	DATABASE EMBL 'Online! HS37245, accession number H93373, 3 December 1995 (1995-12-03) L. HILLIER ET AL: "The WashU-Merck EST project: yv05a10.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:241818 3' similar to gb:M82968 acrosomal protein SP-10 precursor" XP002183455 abstract & UNPUBLISHED, ---	1-3
A	DATABASE EMBL 'Online! S65606, accession number S65606, 11 November 1993 (1993-11-11) R.M. WRIGHT ET AL: "SP-10= intra-acrosomal protein" XP002183456 abstract & R.M. WRIGHT ET AL: "Clonin and characterization of the gene coding for the human acrosomal protein SP-10" BIOL. REPROD., vol. 49, no. 2, 1993, pages 316-325, ---	1-3
P,X	DATABASE EMBL 'Online! Accession number AX086664, 6 March 2001 (2001-03-06) S. WIEMANN: "german human Genome project. Sequence 616 from patent wo0112659" XP002183457 abstract & WO 01 12659 A (GERMAN HUMAN GENOME PROJECT) 22 February 2001 (2001-02-22) ---	1-17, 19-24
	-/--	



## FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

Continuation of Box I.2

Claims Nos.: 30, 31, 44-45, 69-70 all partially

Claim 30, 41 and 45 refer to a selective binding agent of the polypeptide, claim 44 refers to a method of treatment using a selective binding agent without giving a true technical characterization. Moreover, no such compounds are defined in the application. In consequence, the scope of said claims is ambiguous and vague, and their subject-matter is not sufficiently disclosed and supported.

No search can be carried out for such purely speculative claims whose wording is, in fact, a mere recitation of the results to be achieved.

But a partial search has been carried out as far as the selective binding agent is an antibody as mentioned in claim 31.

Claim 69 refers to an antagonist of iL-17 receptor like polypeptide activity and claim 70 refers to a method of reducing cellular production of iL-17 receptor-like polypeptide using a nucleic acid encoding an antagonist of iL-17 receptor like polypeptide without giving a true technical characterization. Moreover, no such compounds are defined in the application. In consequence, the scope of said claims is ambiguous and vague, and their subject-matter is not sufficiently disclosed and supported.

No search can be carried out for such purely speculative claims whose wording is, in fact, a mere recitation of the results to be achieved.

But a partial search has been carried out as far as the antagonist is an antibody against iL-17 receptor like polypeptide or an antisense oligonucleotide.

The applicant's attention is drawn to the fact that claims, or parts of claims, relating to inventions in respect of which no international search report has been established need not be the subject of an international preliminary examination (Rule 66.1(e) PCT). The applicant is advised that the EPO policy when acting as an International Preliminary Examining Authority is normally not to carry out a preliminary examination on matter which has not been searched. This is the case irrespective of whether or not the claims are amended following receipt of the search report or during any Chapter II procedure.

